

SEQUENCE LISTING

<110> F. Hoffmann-La Roche AG
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30 Leu Lys Lys Ser Asn Ala Pro Leu Val Asn Val Thr Leu Tyr Tyr Glu
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25 Pro Asp Ile Tyr Leu Arg Asp Leu Lys Val Glu Asp Ile Pro Leu
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30 Ala Arg Ile Thr Leu Pro Asp Phe Arg Leu Pro Glu Ile Ala Ile
 2705 2710 2715

35 Pro Glu Phe Ile Ile Pro Thr Leu Asn Leu Asn Asp Phe Gln Val
 2720 2725 2730

40 Pro Asp Leu His Ile Pro Glu Phe Gln Leu Pro His Ile Ser His
 2735 2740 2745

45 Thr Ile Glu Val Pro Thr Phe Gly Lys Leu Tyr Ser Ile Leu Lys
 2750 2755 2760

50 Ile Gln Ser Pro Leu Phe Thr Leu Asp Ala Asn Ala Asp Ile Gly
 2765 2770 2775

55 Asn Gly Thr Thr Ser Ala Asn Glu Ala Gly Ile Ala Ala Ser Ile
 2780 2785 2790

60 Thr Ala Lys Gly Glu Ser Lys Leu Glu Val Leu Asn Phe Asp Phe
 2795 2800 2805

65 Gln Ala Asn Ala Gln Leu Ser Asn Pro Lys Ile Asn Pro Leu Ala
 2810 2815 2820

70 Leu Lys Glu Ser Val Lys Phe Ser Ser Lys Tyr Leu Arg Thr Glu
 2825 2830 2835

His Gly Ser Glu Met Leu Phe Phe Gly Asn Ala Ile Glu Gly Lys
 2840 2845 2850

75 Ser Asn Thr Val Ala Ser Leu His Thr Glu Lys Asn Thr Leu Glu
 2855 2860 2865

80 Leu Ser Asn Gly Val Ile Val Lys Ile Asn Asn Gln Leu Thr Leu

	2870	2875	2880
5	Asp Ser Asn Thr Lys Tyr Phe His Lys Leu Asn Ile Pro Lys Leu 2885 2890 2895		
10	Asp Phe Ser Ser Gln Ala Asp Leu Arg Asn Glu Ile Lys Thr Leu 2900 2905 2910		
15	Leu Lys Ala Gly His Ile Ala Trp Thr Ser Ser Gly Lys Gly Ser 2915 2920 2925		
20	Trp Lys Trp Ala Cys Pro Arg Phe Ser Asp Glu Gly Thr His Glu 2930 2935 2940		
25	Ser Gln Ile Ser Phe Thr Ile Glu Gly Pro Leu Thr Ser Phe Gly 2945 2950 2955		
30	Leu Ser Asn Lys Ile Asn Ser Lys His Leu Arg Val Asn Gln Asn 2960 2965 2970		
35	Leu Val Tyr Glu Ser Gly Ser Leu Asn Phe Ser Lys Leu Glu Ile 2975 2980 2985		
40	Gln Ser Gln Val Asp Ser Gln His Val Gly His Ser Val Leu Thr 2990 2995 3000		
45	Ala Lys Gly Met Ala Leu Phe Gly Glu Gly Lys Ala Glu Phe Thr 3005 3010 3015		
50	Gly Arg His Asp Ala His Leu Asn Gly Lys Val Ile Gly Thr Leu 3020 3025 3030		
55	Lys Asn Ser Leu Phe Phe Ser Ala Gln Pro Phe Glu Ile Thr Ala 3035 3040 3045		
60	Ser Thr Asn Asn Glu Gly Asn Leu Lys Val Arg Phe Pro Leu Arg 3050 3055 3060		
65	Leu Thr Gly Lys Ile Asp Phe Leu Asn Asn Tyr Ala Leu Phe Leu 3065 3070 3075		
70	Ser Pro Ser Ala Gln Gln Ala Ser Trp Gln Val Ser Ala Arg Phe 3080 3085 3090		
	Asn Gln Tyr Lys Tyr Asn Gln Asn Phe Ser Ala Gly Asn Asn Glu 3095 3100 3105		
	Asn Ile Met Glu Ala His Val Gly Ile Asn Gly Glu Ala Asn Leu 3110 3115 3120		
	Asp Phe Leu Asn Ile Pro Leu Thr Ile Pro Glu Met Arg Leu Pro 3125 3130 3135		

Tyr Thr Ile Ile Thr Thr Pro Pro Leu Lys Asp Phe Ser Leu Trp
 3140 3145 3150

5 Glu Lys Thr Gly Leu Lys Glu Phe Leu Lys Thr Thr Lys Gln Ser
 3155 3160 3165

10 Phe Asp Leu Ser Val Lys Ala Gln Tyr Lys Lys Asn Lys His Arg
 3170 3175 3180

15 His Ser Ile Thr Asn Pro Leu Ala Val Leu Cys Glu Phe Ile Ser
 3185 3190 3195

20 Gln Ser Ile Lys Ser Phe Asp Arg His Phe Glu Lys Asn Arg Asn
 3200 3205 3210

25 Asn Ala Leu Asp Phe Val Thr Lys Ser Tyr Asn Glu Thr Lys Ile
 3215 3220 3225

30 Lys Phe Asp Lys Tyr Lys Ala Glu Lys Ser His Asp Glu Leu Pro
 3230 3235 3240

35 Arg Thr Phe Gln Ile Pro Gly Tyr Thr Val Pro Val Val Asn Val
 3245 3250 3255

40 Glu Val Ser Pro Phe Thr Ile Glu Met Ser Ala Phe Gly Tyr Val
 3260 3265 3270

45 Phe Pro Lys Ala Val Ser Met Pro Ser Phe Ser Ile Leu Gly Ser
 3275 3280 3285

50 Asp Val Arg Val Pro Ser Tyr Thr Leu Ile Leu Pro Ser Leu Glu
 3290 3295 3300

55 Leu Pro Val Leu His Val Pro Arg Asn Leu Lys Leu Ser Leu Pro
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60 His Phe Lys Glu Leu Cys Thr Ile Ser His Ile Phe Ile Pro Ala
 3320 3325 3330

65 Met Gly Asn Ile Thr Tyr Asp Phe Ser Phe Lys Ser Val Ile
 3335 3340 3345

70 Thr Leu Asn Thr Asn Ala Glu Leu Phe Asn Gln Ser Asp Ile Val
 3350 3355 3360

Ala His Leu Leu Ser Ser Ser Ser Ser Val Ile Asp Ala Leu Gln
 3365 3370 3375

Tyr Lys Leu Glu Gly Thr Thr Arg Leu Thr Arg Lys Arg Gly Leu
 3380 3385 3390

75 Lys Leu Ala Thr Ala Leu Ser Leu Ser Asn Lys Phe Val Glu Gly

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10	Ser Val Ala Lys Thr Thr	Lys 3430	Ala Glu Ile Pro Ile	Leu Arg Met
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	3455		3465	
25	Leu Tyr Ser Thr Ala Lys	Gly 3475	Ala Val Asp His Lys	Leu Ser Leu
	3470		3480	
30	Glu Ser Leu Thr Ser Tyr	Phe 3490	Ser Ile Glu Ser Ser	Thr Lys Gly
	3485		3495	
35	Asp Val Lys Gly Ser Val	Leu 3505	Ser Arg Glu Tyr Ser	Gly Thr Ile
	3500		3510	
40	Ala Ser Glu Ala Asn Thr	Tyr 3520	Leu Asn Ser Lys Ser	Thr Arg Ser
	3515		3525	
45	Ser Val Lys Leu Gln Gly	Thr 3535	Ser Lys Ile Asp Asp	Ile Trp Asn
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50	Leu Glu Val Lys Glu Asn	Phe 3550	Ala Gly Glu Ala Thr	Leu Gln Arg
	3545		3555	
55	Ile Tyr Ser Leu Trp Glu	His 3565	Ser Thr Lys Asn His	Leu Gln Leu
	3560		3570	
60	Glu Gly Leu Phe Phe Thr	Asn 3580	Gly Glu His Thr Ser	Lys Ala Thr
	3575		3585	
65	Leu Glu Leu Ser Pro Trp	Gln 3595	Met Ser Ala Leu Val	Gln Val His
	3590		3600	
70	Ala Ser Gln Pro Ser Ser	Phe 3610	His Asp Phe Pro Asp	Leu Gly Gln
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	3620		3630	
80	Lys Asn Glu Val Arg Ile	His 3640	Ser Gly Ser Phe Gln	Ser Gln Val
	3635		3645	
85	Glu Leu Ser Asn Asp Gln	Glu 3655	Lys Ala His Leu Asp	Ile Ala Gly
	3650		3660	

Ser Leu Glu Gly His Leu Arg Phe Leu Lys Asn Ile Ile Leu Pro
 3665 3670 3675
 5 Val Tyr Asp Lys Ser Leu Trp Asp Phe Leu Lys Leu Asp Val Thr
 3680 3685 3690
 10 Thr Ser Ile Gly Arg Arg Gln His Leu Arg Val Ser Thr Ala Phe
 3695 3700 3705
 15 Val Tyr Thr Lys Asn Pro Asn Gly Tyr Ser Phe Ser Ile Pro Val
 3710 3715 3720
 20 Lys Val Leu Ala Asp Lys Phe Ile Thr Pro Gly Leu Lys Leu Asn
 3725 3730 3735
 25 Asp Leu Asn Ser Val Leu Val Met Pro Thr Phe His Val Pro Phe
 3740 3745 3750
 30 Thr Asp Leu Gln Val Pro Ser Cys Lys Leu Asp Phe Arg Glu Ile
 3755 3760 3765
 35 Gln Ile Tyr Lys Lys Leu Arg Thr Ser Ser Phe Ala Leu Asn Leu
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 40 Pro Thr Leu Pro Glu Val Lys Phe Pro Glu Val Asp Val Leu Thr
 3785 3790 3795
 Lys Tyr Ser Gln Pro Glu Asp Ser Leu Ile Pro Phe Glu Ile
 3800 3805 3810
 45 Thr Val Pro Glu Ser Gln Leu Thr Val Ser Gln Phe Thr Leu Pro
 3815 3820 3825
 50 Lys Ser Val Ser Asp Gly Ile Ala Ala Leu Asp Leu Asn Ala Val
 3830 3835 3840
 55 Ala Asn Lys Ile Ala Asp Phe Glu Leu Pro Thr Ile Ile Val Pro
 3845 3850 3855
 Glu Gln Thr Ile Glu Ile Pro Ser Ile Lys Phe Ser Val Pro Ala
 3860 3865 3870
 60 Gly Ile Val Ile Pro Ser Phe Gln Ala Leu Thr Ala Arg Phe Glu
 3875 3880 3885
 65 Val Asp Ser Pro Val Tyr Asn Ala Thr Trp Ser Ala Ser Leu Lys
 3890 3895 3900
 Asn Lys Ala Asp Tyr Val Glu Thr Val Leu Asp Ser Thr Cys Ser
 3905 3910 3915
 70 Ser Thr Val Gln Phe Leu Glu Tyr Glu Leu Asn Val Leu Gly Thr

	3920	3925	3930
5	His Lys Ile Glu Asp Gly Thr 3935	Leu Ala Ser Lys Thr 3940	Lys Gly Thr 3945
10	Leu Ala His Arg Asp Phe Ser 3950	Ala Glu Tyr Glu 3955	Glu Asp Gly Lys 3960
15	Phe Glu Gly Leu Gln Glu Trp 3965	Glu Gly Lys Ala His 3970	Leu Asn Ile 3975
20	Lys Ser Pro Ala Phe Thr Asp 3980	Leu His Leu Arg Tyr 3985	Gln Lys Asp 3990
25	Lys Lys Gly Ile Ser Thr Ser 3995	Ala Ala Ser Pro Ala Val 4000	Gly Thr 4005
30	Val Gly Met Asp Met Asp Glu 4010	Asp Asp Asp Phe Ser 4015	Lys Trp Asn 4020
35	Phe Tyr Tyr Ser Pro Gln Ser 4025	Ser Pro Asp Lys Lys 4030	Leu Thr Ile 4035
40	Phe Lys Thr Glu Leu Arg Val 4040	Arg Glu Ser Asp Glu 4045	Glu Thr Gln 4050
45	Ile Lys Val Asn Trp Glu Glu 4055	Glu Ala Ala Ser Gly 4060	Leu Leu Thr 4065
50	Ser Leu Lys Asp Asn Val Pro 4070	Lys Ala Thr Gly Val 4075	Leu Tyr Asp 4080
55	Tyr Val Asn Lys Tyr His Trp 4085	Glu His Thr Gly Leu 4090	Thr Leu Arg 4095
60	Glu Val Ser Ser Lys Leu Arg 4100	Arg Asn Leu Gln Asn 4105	Asn Ala Glu 4110
65	Trp Val Tyr Gln Gly Ala Ile 4115	Arg Gln Ile Asp Asp 4120	Ile Asp Val 4125
70	Arg Phe Gln Lys Ala Ala Ser 4130	Gly Thr Thr Gly Thr 4135	Tyr Gln Glu 4140
	Trp Lys Asp Lys Ala Gln Asn 4145	Leu Tyr Gln Glu Leu 4150	Leu Thr Gln 4155
	Glu Gly Gln Ala Ser Phe Gln 4160	Gly Leu Lys Asp Asn 4165	Val Phe Asp 4170
	Gly Leu Val Arg Val Thr Gln 4175	Lys Phe His Met Lys 4180	Val Lys His 4185

Leu Ile Asp Ser Leu Ile Asp Phe Leu Asn Phe Pro Arg Phe Gln
4190 4195 4200

5 Phe Pro Gly Lys Pro Gly Ile Tyr Thr Arg Glu Glu Leu Cys Thr
4205 4210 4215

10 Met Phe Ile Arg Glu Val Gly Thr Val Leu Ser Gln Val Tyr Ser
4220 4225 4230

15 Lys Val His Asn Gly Ser Glu Ile Leu Phe Ser Tyr Phe Gln Asp
4235 4240 4245

20 Leu Val Ile Thr Leu Pro Phe Glu Leu Arg Lys His Lys Leu Ile
4250 4255 4260

25 Asp Val Ile Ser Met Tyr Arg Glu Leu Leu Lys Asp Leu Ser Lys
4265 4270 4275

30 Glu Ala Gln Glu Val Phe Lys Ala Ile Gln Ser Leu Lys Thr Thr
4280 4285 4290

35 Glu Val Leu Arg Asn Leu Gln Asp Leu Leu Gln Phe Ile Phe Gln
4295 4300 4305

40 Leu Ile Glu Asp Asn Ile Lys Gln Leu Lys Glu Met Lys Phe Thr
4310 4315 4320

45 Tyr Leu Ile Asn Tyr Ile Gln Asp Glu Ile Asn Thr Ile Phe Asn
4325 4330 4335

50 Asp Tyr Ile Pro Tyr Val Phe Lys Leu Leu Lys Glu Asn Leu Cys
4340 4345 4350

55 Leu Asn Leu His Lys Phe Asn Glu Phe Ile Gln Asn Glu Leu Gln
4355 4360 4365

60 Glu Ala Ser Gln Glu Leu Gln Gln Ile His Gln Tyr Ile Met Ala
4370 4375 4380

65 Leu Arg Glu Glu Tyr Phe Asp Pro Ser Ile Val Gly Trp Thr Val
4385 4390 4395

70 Lys Tyr Tyr Glu Leu Glu Lys Ile Val Ser Leu Ile Lys Asn
4400 4405 4410

75 Leu Leu Val Ala Leu Lys Asp Phe His Ser Glu Tyr Ile Val Ser
4415 4420 4425

80 Ala Ser Asn Phe Thr Ser Gln Leu Ser Ser Gln Val Glu Gln Phe
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85 Leu His Arg Asn Ile Gln Glu Tyr Leu Ser Ile Leu Thr Asp Pro

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15	Ser Asp Tyr His Gln Gln Phe Arg Tyr Lys Leu Gln Asp Phe Ser 4490 4495 4500		
20	Asp Gln Leu Ser Asp Tyr Tyr Glu Lys Phe Ile Ala Glu Ser Lys 4505 4510 4515		
25	Arg Leu Ile Asp Leu Ser Ile Gln Asn Tyr His Thr Phe Leu Ile 4520 4525 4530		
30	Tyr Ile Thr Glu Leu Leu Lys Lys Leu Gln Ser Thr Thr Val Met 4535 4540 4545		
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55	Ile Asp Ile Tyr Ser Leu Thr Val Asp Ser Arg Val Ser Ser Arg Phe 35 40 45		
60	Ala His Thr Val Val Thr Ser Arg Val Val Asn Arg Ala Asn Thr Val 50 55 60		
65	Gln Glu Ala Thr Phe Gln Met Glu Leu Pro Lys Lys Ala Phe Ile Thr 65 70 75 80		
70	Asn Phe Ser Met Asn Ile Asp Gly Met Thr Tyr Pro Gly Ile Ile Lys 85 90 95		
	Glu Lys Ala Glu Ala Gln Ala Gln Tyr Ser Ala Ala Val Ala Lys Gly 100 105 110		

Lys Ser Ala Gly Leu Val Lys Ala Thr Gly Arg Asn Met Glu Gln Phe
115 120 125

5 Gln Val Ser Val Ser Val Ala Pro Asn Ala Lys Ile Thr Phe Glu Leu
130 135 140

10 Val Tyr Glu Glu Leu Leu Lys Arg Arg Leu Gly Val Tyr Glu Leu Leu
145 150 155 160

15 Leu Lys Val Arg Pro Gln Gln Leu Val Lys His Leu Gln Met Asp Ile
165 170 175

20 His Ile Phe Glu Pro Gln Gly Ile Ser Phe Leu Glu Thr Glu Ser Thr
180 185 190

25 Phe Met Thr Asn Gln Leu Val Asp Ala Leu Thr Thr Trp Gln Asn Lys
195 200 205

30 Thr Lys Ala His Ile Arg Phe Lys Pro Thr Leu Ser Gln Gln Gln Lys
210 215 220

35 Ser Pro Glu Gln Gln Glu Thr Val Leu Asp Gly Asn Leu Ile Ile Arg
225 230 235 240

40 Tyr Asp Val Asp Arg Ala Ile Ser Gly Gly Ser Ile Gln Ile Glu Asn
245 250 255

45 Gly Tyr Phe Val His Tyr Phe Ala Pro Glu Gly Leu Thr Thr Met Pro
260 265 270

50 Lys Asn Val Val Phe Val Ile Asp Lys Ser Gly Ser Met Ser Gly Arg
275 280 285

55 Lys Ile Gln Gln Thr Arg Glu Ala Leu Ile Lys Ile Leu Asp Asp Leu
290 295 300

60 Ser Pro Arg Asp Gln Phe Asn Leu Ile Val Phe Ser Thr Glu Ala Thr
305 310 315 320

65 Gln Trp Arg Pro Ser Leu Val Pro Ala Ser Ala Glu Asn Val Asn Lys
325 330 335

70 Ala Arg Ser Phe Ala Ala Gly Ile Gln Ala Leu Gly Gly Thr Asn Ile
340 345 350

75 Asn Asp Ala Met Leu Met Ala Val Gln Leu Leu Asp Ser Ser Asn Gln
355 360 365

80 Glu Glu Arg Leu Pro Glu Gly Ser Val Ser Leu Ile Ile Leu Leu Thr
370 375 380

85 Asp Gly Asp Pro Thr Val Gly Glu Thr Asn Pro Arg Ser Ile Gln Asn

385 390 395 400

5 Asn Val Arg Glu Ala Val Ser Gly Arg Tyr Ser Leu Phe Cys Leu Gly
405 410 415

Phe Gly Phe Asp Val Ser Tyr Ala Phe Leu Glu Lys Leu Ala Leu Asp
420 425 430

10 Asn Gly Gly Leu Ala Arg Arg Ile His Glu Asp Ser Asp Ser Ala Leu
435 440 445

15 Gln Leu Gln Asp Phe Tyr Gln Glu Val Ala Asn Pro Leu Leu Thr Ala
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20 Val Thr Phe Glu Tyr Pro Ser Asn Ala Val Glu Glu Val Thr Gln Asn
465 470 475 480

25 Asn Phe Arg Leu Leu Phe Lys Gly Ser Glu Met Val Val Ala Gly Lys
485 490 495

30 Leu Gln Asp Arg Gly Pro Asp Val Leu Thr Ala Thr Val Ser Gly Lys
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Leu Pro Thr Gln Asn Ile Thr Phe Gln Thr Glu Ser Ser Val Ala Glu
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35 Gln Glu Ala Glu Phe Gln Ser Pro Lys Tyr Ile Phe His Asn Phe Met
530 535 540

40 Glu Arg Leu Trp Ala Tyr Leu Thr Ile Gln Gln Leu Leu Glu Gln Thr
545 550 555 560

45 Val Ser Ala Ser Asp Ala Asp Gln Gln Ala Leu Arg Asn Gln Ala Leu
565 570 575

50 Asn Leu Ser Leu Ala Tyr Ser Phe Val Thr Pro Leu Thr Ser Met Val
580 585 590

Val Thr Lys Pro Asp Asp Gln Glu Gln Ser Gln Val Ala Glu Lys Pro
595 600 605

55 Met Glu Gly Glu Ser Arg Asn Arg Asn Val His Ser Gly Ser Thr Phe
610 615 620

60 Phe Lys Tyr Tyr Leu Gln Gly Ala Lys Ile Pro Lys Pro Glu Ala Ser
625 630 635 640

65 Phe Ser Pro Arg Arg Gly Trp Asn Arg Gln Ala Gly Ala Ala Gly Ser
645 650 655

Arg Met Asn Phe Arg Pro Gly Val Leu Ser Ser Arg Gln Leu Gly Leu
660 665 670

Pro Gly Pro Pro Asp Val Pro Asp His Ala Ala Tyr His Pro Phe Arg
675 680 685

5 Arg Leu Ala Ile Leu Pro Ala Ser Ala Pro Pro Ala Thr Ser Asn Pro
690 695 700

10 Asp Pro Ala Val Ser Arg Val Met Asn Met Lys Ile Glu Glu Thr Thr
705 710 715 720

15 Met Thr Thr Gln Thr Pro Ala Pro Ile Gln Ala Pro Ser Ala Ile Leu
725 730 735

20 Pro Leu Pro Gly Gln Ser Val Glu Arg Leu Cys Val Asp Pro Arg His
740 745 750

Arg Gln Gly Pro Val Asn Leu Leu Ser Asp Pro Glu Gln Gly Val Glu
755 760 765

25 Val Thr Gly Gln Tyr Glu Arg Glu Lys Ala Gly Phe Ser Trp Ile Glu
770 775 780

30 Val Thr Phe Lys Asn Pro Leu Val Trp Val His Ala Ser Pro Glu His
785 790 795 800

35 Val Val Val Thr Arg Asn Arg Arg Ser Ser Ala Tyr Lys Trp Lys Glu
805 810 815

40 Thr Leu Phe Ser Val Met Pro Gly Leu Lys Met Thr Met Asp Lys Thr
820 825 830

Gly Leu Leu Leu Ser Asp Pro Asp Lys Val Thr Ile Gly Leu Leu
835 840 845

45 Phe Trp Asp Gly Arg Gly Glu Gly Leu Arg Leu Leu Leu Arg Asp Thr
850 855 860

50 Asp Arg Phe Ser Ser His Val Gly Gly Thr Leu Gly Gln Phe Tyr Gln
865 870 875 880

55 Glu Val Leu Trp Gly Ser Pro Ala Ala Ser Asp Asp Gly Arg Arg Thr
885 890 895

60 Leu Arg Val Gln Gly Asn Asp His Ser Ala Thr Arg Glu Arg Arg Leu
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Asp Tyr Gln Glu Gly Pro Pro Gly Val Glu Ile Ser Cys Trp Ser Val
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65 Glu Leu
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<309> 1986-07-21

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Leu Gly Val Pro Leu Ser Val Gly Val Gln Leu Gln Asp Val Pro Arg
35 40 45

25

Gly Gln Val Val Lys Gly Ser Val Phe Leu Arg Asn Pro Ser Arg Asn
50 55 60

30

Asn Val Pro Cys Ser Pro Lys Val Asp Phe Thr Leu Ser Ser Glu Arg
65 70 75 80

Asp Phe Ala Leu Leu Ser Leu Gln Val Pro Leu Lys Asp Ala Lys Ser
85 90 95

35

Cys Gly Leu His Gln Leu Leu Arg Gly Pro Glu Val Gln Leu Val Ala
100 105 110

40

His Ser Pro Trp Leu Lys Asp Ser Leu Ser Arg Thr Thr Asn Ile Gln
115 120 125

45

Gly Ile Asn Leu Leu Phe Ser Ser Arg Arg Gly His Leu Phe Leu Gln
130 135 140

50

Thr Asp Gln Pro Ile Tyr Asn Pro Gly Gln Arg Val Arg Tyr Arg Val
145 150 155 160

Phe Ala Leu Asp Gln Lys Met Arg Pro Ser Thr Asp Thr Ile Thr Val
165 170 175

55

Met Val Glu Asn Ser His Gly Leu Arg Val Arg Lys Lys Glu Val Tyr
180 185 190

60

Met Pro Ser Ser Ile Phe Gln Asp Asp Phe Val Ile Pro Asp Ile Ser
195 200 205

65

Glu Pro Gly Thr Trp Lys Ile Ser Ala Arg Phe Ser Asp Gly Leu Glu
210 215 220

70

Ser Asn Ser Ser Thr Gln Phe Glu Val Lys Lys Tyr Val Leu Pro Asn
225 230 235 240

Phe Glu Val Lys Ile Thr Pro Gly Lys Pro Tyr Ile Leu Thr Val Pro
245 250 255

5 Gly His Leu Asp Glu Met Gln Leu Asp Ile Gln Ala Arg Tyr Ile Tyr
260 265 270

10 Gly Lys Pro Val Gln Gly Val Ala Tyr Val Arg Phe Gly Leu Leu Asp
275 280 285

15 Glu Asp Gly Lys Lys Thr Phe Phe Arg Gly Leu Glu Ser Gln Thr Lys
290 295 300

20 Leu Val Asn Gly Gln Ser His Ile Ser Leu Ser Lys Ala Glu Phe Gln
305 310 315 320

Asp Ala Leu Glu Lys Leu Asn Met Gly Ile Thr Asp Leu Gln Gly Leu
325 330 335

25 Arg Leu Tyr Val Ala Ala Ala Ile Ile Glu Ser Pro Gly Gly Glu Met
340 345 350

30 Glu Glu Ala Glu Leu Thr Ser Trp Tyr Phe Val Ser Ser Pro Phe Ser
355 360 365

35 Leu Asp Leu Ser Lys Thr Lys Arg His Leu Val Pro Gly Ala Pro Phe
370 375 380

Leu Leu Gln Ala Leu Val Arg Glu Met Ser Gly Ser Pro Ala Ser Gly
385 390 395 400

40 Ile Pro Val Lys Val Ser Ala Thr Val Ser Ser Pro Gly Ser Val Pro
405 410 415

45 Glu Val Gln Asp Ile Gln Gln Asn Thr Asp Gly Ser Gly Gln Val Ser
420 425 430

50 Ile Pro Ile Ile Pro Gln Thr Ile Ser Glu Leu Gln Leu Ser Val
435 440 445

55 Ser Ala Gly Ser Pro His Pro Ala Ile Ala Arg Leu Thr Val Ala Ala
450 455 460

Pro Pro Ser Gly Gly Pro Gly Phe Leu Ser Ile Glu Arg Pro Asp Ser
465 470 475 480

60 Arg Pro Pro Arg Val Gly Asp Thr Leu Asn Leu Asn Leu Arg Ala Val
485 490 495

65 Gly Ser Gly Ala Thr Phe Ser His Tyr Tyr Tyr Met Ile Leu Ser Arg
500 505 510

70 Gly Gln Ile Val Phe Met Asn Arg Glu Pro Lys Arg Thr Leu Thr Ser

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20	Gly Ala Lys Gln Tyr Arg Asn Gly Glu Ser Val Lys Leu His Leu Glu 580 585 590		
25	Thr Asp Ser Leu Ala Leu Val Ala Leu Gly Ala Leu Asp Thr Ala Leu 595 600 605		
30	Tyr Ala Ala Gly Ser Lys Ser His Lys Pro Leu Asn Met Gly Lys Val 610 615 620		
35	Phe Glu Ala Met Asn Ser Tyr Asp Leu Gly Cys Gly Pro Gly Gly Gly 625 630 635 640		
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45	Gly Asp Gln Trp Thr Leu Ser Arg Lys Arg Leu Ser Cys Pro Lys Glu 660 665 670		
50	Lys Thr Thr Arg Lys Lys Arg Asn Val Asn Phe Gln Lys Ala Ile Asn 675 680 685		
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65	Ala Arg Val Gln Gln Pro Asp Cys Arg Glu Pro Phe Leu Ser Cys Cys 725 730 735		
70	Gln Phe Ala Glu Ser Leu Arg Lys Lys Ser Arg Asp Lys Gly Gln Ala 740 745 750		
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80	Glu Asp Asp Ile Pro Val Arg Ser Phe Phe Pro Glu Asn Trp Leu Trp 770 775 780		
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Asp Ser Leu Thr Thr Trp Glu Ile His Gly Leu Ser Leu Ser Lys Thr
805 810 815

5 Lys Gly Leu Cys Val Ala Thr Pro Val Gln Leu Arg Val Phe Arg Glu
820 825 830

10 Phe His Leu His Leu Arg Leu Pro Met Ser Val Arg Arg Phe Glu Gln
835 840 845

15 Leu Glu Leu Arg Pro Val Leu Tyr Asn Tyr Leu Asp Lys Asn Leu Thr
850 855 860

20 Val Ser Val His Val Ser Pro Val Glu Gly Leu Cys Leu Ala Gly Gly
865 870 875 880

Gly Gly Leu Ala Gln Gln Val Leu Val Pro Ala Gly Ser Ala Arg Pro
885 890 895

25 Val Ala Phe Ser Val Val Pro Thr Ala Ala Ala Ala Val Ser Leu Lys
900 905 910

30 Val Val Ala Arg Gly Ser Phe Glu Phe Pro Val Gly Asp Ala Val Ser
915 920 925

35 Lys Val Leu Gln Ile Glu Lys Glu Gly Ala Ile His Arg Glu Glu Leu
930 935 940

40 Val Tyr Glu Leu Asn Pro Leu Asp His Arg Gly Arg Thr Leu Glu Ile
945 950 955 960

45 Pro Gly Asn Ser Asp Pro Asn Met Ile Pro Asp Gly Asp Phe Asn Ser
965 970 975

Tyr Val Arg Val Thr Ala Ser Asp Pro Leu Asp Thr Leu Gly Ser Glu
980 985 990

50 Gly Ala Leu Ser Pro Gly Gly Val Ala Ser Leu Leu Arg Leu Pro Arg
995 1000 1005

55 Gly Cys Gly Glu Gln Thr Met Ile Tyr Leu Ala Pro Thr Leu Ala
1010 1015 1020

60 Ala Ser Arg Tyr Leu Asp Lys Thr Glu Gln Trp Ser Thr Leu Pro
1025 1030 1035

65 Pro Glu Thr Lys Asp His Ala Val Asp Leu Ile Gln Lys Gly Tyr
1040 1045 1050

70 Met Arg Ile Gln Gln Phe Arg Lys Ala Asp Gly Ser Tyr Ala Ala
1055 1060 1065

Trp Leu Ser Arg Asp Ser Ser Thr Trp Leu Thr Ala Phe Val Leu

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10	Lys Leu 1100	Gln Glu Thr Ser Asn 1105	Trp Leu Leu Ser Gln 1110	Gln Gln Ala
15	Asp Gly 1115	Ser Phe Gln Asp Pro 1120	Cys Pro Val Leu Asp 1125	Arg Ser Met
20	Gln Gly 1130	Gly Leu Val Gly Asn 1135	Asp Glu Thr Val Ala 1140	Leu Thr Ala
25	Phe Val 1145	Thr Ile Ala Leu His 1150	His Gly Leu Ala Val 1155	Phe Gln Asp
30	Glu Gly 1160	Ala Glu Pro Leu Lys 1165	Gln Arg Val Glu Ala 1170	Ser Ile Ser
35	Lys Ala 1175	Asn Ser Phe Leu Gly 1180	Glu Lys Ala Ser Ala 1185	Gly Leu Leu
40	Gly Ala 1190	His Ala Ala Ala Ile 1195	Thr Ala Tyr Ala Leu 1200	Ser Leu Thr
45	Lys Ala 1205	Pro Val Asp Leu Leu 1210	Gly Val Ala His Asn 1215	Asn Leu Met
50	Ala Met 1220	Ala Gln Glu Thr Gly 1225	Asp Asn Leu Tyr Trp 1230	Gly Ser Val
55	Thr Gly 1235	Ser Gln Ser Asn Ala 1240	Val Ser Pro Thr Pro 1245	Ala Pro Arg
60	Asn Pro 1250	Ser Asp Pro Met Pro 1255	Gln Ala Pro Ala Leu 1260	Trp Ile Glu
65	Thr Thr 1265	Ala Tyr Ala Leu Leu 1270	His Leu Leu Leu His 1275	Glu Gly Lys
70	Ala Glu 1280	Met Ala Asp Gln Ala 1285	Ser Ala Trp Leu Thr 1290	Arg Gln Gly
	Ser Phe 1295	Gln Gly Gly Phe Arg 1300	Ser Thr Gln Asp Thr 1305	Val Ile Ala
	Leu Asp 1310	Ala Leu Ser Ala Tyr 1315	Trp Ile Ala Ser His 1320	Thr Thr Glu
	Glu Arg 1325	Gly Leu Asn Val Thr 1330	Leu Ser Ser Thr Gly 1335	Arg Asn Gly

Phe Lys Ser His Ala Leu Gln Leu Asn Asn Arg Gln Ile Arg Gly
 1340 1345 1350
 5 Leu Glu Glu Glu Leu Gln Phe Ser Leu Gly Ser Lys Ile Asn Val
 1355 1360 1365
 10 Lys Val Gly Gly Asn Ser Lys Gly Thr Leu Lys Val Leu Arg Thr
 1370 1375 1380
 15 Tyr Asn Val Leu Asp Met Lys Asn Thr Thr Cys Gln Asp Leu Gln
 1385 1390 1395
 20 Ile Glu Val Thr Val Lys Gly His Val Glu Tyr Thr Met Glu Ala
 1400 1405 1410
 Asn Glu Asp Tyr Glu Asp Tyr Glu Tyr Asp Glu Leu Pro Ala Lys
 1415 1420 1425
 25 Asp Asp Pro Asp Ala Pro Leu Gln Pro Val Thr Pro Leu Gln Leu
 1430 1435 1440
 30 Phe Glu Gly Arg Arg Asn Arg Arg Arg Glu Ala Pro Lys Val
 1445 1450 1455
 35 Val Glu Glu Gln Glu Ser Arg Val His Tyr Thr Val Cys Ile Trp
 1460 1465 1470
 40 Arg Asn Gly Lys Val Gly Leu Ser Gly Met Ala Ile Ala Asp Val
 1475 1480 1485
 Thr Leu Leu Ser Gly Phe His Ala Leu Arg Ala Asp Leu Glu Lys
 1490 1495 1500
 45 Leu Thr Ser Leu Ser Asp Arg Tyr Val Ser His Phe Glu Thr Glu
 1505 1510 1515
 50 Gly Pro His Val Leu Leu Tyr Phe Asp Ser Val Pro Thr Ser Arg
 1520 1525 1530
 55 Glu Cys Val Gly Phe Glu Ala Val Gln Glu Val Pro Val Gly Leu
 1535 1540 1545
 60 Val Gln Pro Ala Ser Ala Thr Leu Tyr Asp Tyr Tyr Asn Pro Glu
 1550 1555 1560
 Arg Arg Cys Ser Val Phe Tyr Gly Ala Pro Ser Lys Ser Arg Leu
 1565 1570 1575
 65 Leu Ala Thr Leu Cys Ser Ala Glu Val Cys Gln Cys Ala Glu Gly
 1580 1585 1590
 70 Lys Cys Pro Arg Gln Arg Arg Ala Leu Glu Arg Gly Leu Gln Asp

	1595	1600	1605
5	Glu Asp Gly Tyr Arg Met Lys	Phe Ala Cys Tyr Tyr	Pro Arg Val
	1610 1615	1620	
10	Glu Tyr Gly Phe Gln Val Lys	Val Leu Arg Glu Asp Ser Arg Ala	
	1625 1630	1635	
15	Ala Phe Arg Leu Phe Glu Thr	Lys Ile Thr Gln Val	Leu His Phe
	1640 1645	1650	
20	Thr Lys Asp Val Lys Ala Ala	Ala Asn Gln Met Arg Asn Phe Leu	
	1655 1660	1665	
25	Val Arg Ala Ser Cys Arg Leu	Arg Leu Glu Pro Gly Lys Glu Tyr	
	1670 1675	1680	
30	Leu Ile Met Gly Leu Asp Gly	Ala Thr Tyr Asp Leu Glu Gly His	
	1685 1690	1695	
35	Pro Gln Tyr Leu Leu Asp Ser	Asn Ser Trp Ile Glu Glu Met Pro	
	1700 1705	1710	
40	Ser Glu Arg Leu Cys Arg Ser	Thr Arg Gln Arg Ala Ala Cys Ala	
	1715 1720	1725	
45	Gln Leu Asn Asp Phe Leu Gln	Glu Tyr Gly Thr Gln Gly Cys Gln	
	1730 1735	1740	
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65	Leu Pro Leu Ala Leu Gly Ser Pro Met Tyr Ser Ile Ile Thr Pro Asn		
	20 25 30		
	Ile Leu Arg Leu Glu Ser Glu Glu Thr Met Val Leu Glu Ala His Asp		
	35 40 45		
70	Ala Gln Gly Asp Val Pro Val Thr Val Thr Val His Asp Phe Pro Gly		
	50 55 60		

Lys Lys Leu Val Leu Ser Ser Glu Lys Thr Val Leu Thr Pro Ala Thr
65 70 75 80

5 Asn His Met Gly Asn Val Thr Phe Thr Ile Pro Ala Asn Arg Glu Phe
85 90 95

10 Lys Ser Glu Lys Gly Arg Asn Lys Phe Val Thr Val Gln Ala Thr Phe
100 105 110

15 Gly Thr Gln Val Val Glu Lys Val Val Leu Val Ser Leu Gln Ser Gly
115 120 125

20 Tyr Leu Phe Ile Gln Thr Asp Lys Thr Ile Tyr Thr Pro Gly Ser Thr
130 135 140

25 Val Leu Tyr Arg Ile Phe Thr Val Asn His Lys Leu Leu Pro Val Gly
145 150 155 160

30 Gln Asp Ser Leu Ser Ser Gln Asn Gln Leu Gly Val Leu Pro Leu Ser
180 185 190

35 Trp Asp Ile Pro Glu Leu Val Asn Met Gly Gln Trp Lys Ile Arg Ala
195 200 205

40 Tyr Tyr Glu Asn Ser Pro Gln Gln Val Phe Ser Thr Glu Phe Glu Val
210 215 220

45 Lys Glu Tyr Val Leu Pro Ser Phe Glu Val Ile Val Glu Pro Thr Glu
225 230 235 240

50 Ala Arg Phe Leu Tyr Gly Lys Lys Val Glu Gly Thr Ala Phe Val Ile
260 265 270

55 Phe Gly Ile Gln Asp Gly Glu Gln Arg Ile Ser Leu Pro Glu Ser Leu
275 280 285

60 Lys Arg Ile Pro Ile Glu Asp Gly Ser Gly Glu Val Val Leu Ser Arg
290 295 300

65 Lys Val Leu Leu Asp Gly Val Gln Asn Leu Arg Ala Glu Asp Leu Val
305 310 315 320

70 Gly Lys Ser Leu Tyr Val Ser Ala Thr Val Ile Leu His Ser Gly Ser
325 330 335

Asp Met Val Gln Ala Glu Arg Ser Gly Ile Pro Ile Val Thr Ser Pro

38

340

345

350

5 - Tyr Gln Ile His Phe Thr Lys Thr Pro Lys Tyr Phe Lys Pro Gly Met
 355 360 365

Pro Phe Asp Leu Met Val Phe Val Thr Asn Pro Asp Gly Ser Pro Ala
370 375 380

10 Tyr Arg Val Pro Val Ala Val Gln Gly Glu Asp Thr Val Gln Ser Leu
385 390 395 400

15 Thr Gln Gly Asp Gly Val Ala Lys Leu Ser Ile Asn Thr His Pro Ser
 405 410 415

20 Gln Lys Pro Leu Ser Ile Thr Val Arg Thr Lys Lys Gln Glu Leu Ser
420 425 430

25 Glu Ala Glu Gln Ala Thr Arg Thr Met Gln Ala Leu Pro Tyr Ser Thr
435 440 445

Val Gly Asn Ser Asn Asn Tyr Leu His Leu Ser Val Leu Arg Thr Glu
450 455 460

Leu Arg Pro Gly Glu Thr Leu Asn Val Asn Phe Leu Leu Arg Met Asp

35 Arg Ala His Glu Ala Lys Ile Arg Tyr Tyr Thr Tyr Leu Ile Met Asn
185 190 195

40 Lys Gly Arg Leu Leu Lys Ala Gly Arg Gln Val Arg Glu Pro Gly Glu

Asp Leu Val Val Leu Pro Leu Ser Ile Thr Thr Asp Phe Ile Pro Ser
515 520 525

Phe Arg Leu Val Ala Tyr Tyr Thr Leu Ile Gly Ala Ser Gly Gln Arg
528 529 530 531 532 533 534 535 536 537 538 539 540

50 Glu Val Val Ala Asp Ser Val Trp Val Asp Val Lys Asp Ser Cys Val

55 Gly Ser Leu Val Val Lys Ser Gly Gln Ser Glu Asp Arg Gln Pro Val

60 Pro Gly Glu Glu Met Thr Leu Lys Ile Glu Gly Asp His Gly Ala Arg

Val Val Leu Val Ala Val Asp Lys Gly Val Phe Val Leu Asn Lys Lys

Asn Lys Leu Thr Gln Ser Lys Ile Trp Asp Val Val Glu Lys Ala Asp

Ile Gly Cys Thr Pro Gly Ser Gly Lys Asp Tyr Ala Gly Val Phe Ser
625 630 635 640

5 Asp Ala Gly Leu Thr Phe Thr Ser Ser Ser Gly Gln Gln Thr Ala Gln
645 650 655

10 Arg Ala Glu Leu Gln Cys Pro Gln Pro Ala Ala Arg Arg Arg Arg Ser
660 665 670

15 Val Gln Leu Thr Glu Lys Arg Met Asp Lys Val Gly Lys Tyr Pro Lys
675 680 685

20 Glu Leu Arg Lys Cys Cys Glu Asp Gly Met Arg Glu Asn Pro Met Arg
690 695 700

25 Phe Ser Cys Gln Arg Arg Thr Arg Phe Ile Ser Leu Gly Glu Ala Cys
705 710 715 720

30 Lys Lys Val Phe Leu Asp Cys Cys Asn Tyr Ile Thr Glu Leu Arg Arg
725 730 735

35 Gln His Ala Arg Ala Ser His Leu Gly Leu Ala Arg Ser Asn Leu Asp
740 745 750

40 Glu Asp Ile Ile Ala Glu Glu Asn Ile Val Ser Arg Ser Glu Phe Pro
755 760 765

45 Glu Ser Trp Leu Trp Asn Val Glu Asp Leu Lys Glu Pro Pro Lys Asn
770 775 780

50 Gly Ile Ser Thr Lys Leu Met Asn Ile Phe Leu Lys Asp Ser Ile Thr
785 790 795 800

55 Thr Trp Glu Ile Leu Ala Val Ser Met Ser Asp Lys Lys Gly Ile Cys
805 810 815

60 Val Ala Asp Pro Phe Glu Val Thr Val Met Gln Asp Phe Phe Ile Asp
820 825 830

65 Leu Arg Leu Pro Tyr Ser Val Val Arg Asn Glu Gln Val Glu Ile Arg
835 840 845

70 Ala Val Leu Tyr Asn Tyr Arg Gln Asn Gln Glu Leu Lys Val Arg Val
850 855 860

75 Glu Leu Leu His Asn Pro Ala Phe Cys Ser Leu Ala Thr Thr Lys Arg
865 870 875 880

80 Arg His Gln Gln Thr Val Thr Ile Pro Pro Lys Ser Ser Leu Ser Val
885 890 895

85 Pro Tyr Val Ile Val Pro Leu Lys Thr Gly Leu Gln Glu Val Glu Val

40

900

905

910

5 Lys Ala Ala Val Tyr His His Phe Ile Ser Asp Gly Val Arg Lys Ser
915 920 925

10 Leu Lys Val Val Pro Glu Gly Ile Arg Met Asn Lys Thr Val Ala Val
930 935 940

Arg Thr Leu Asp Pro Glu Arg Leu Gly Arg Glu Gly Val Gln Lys Glu
945 950 955 960

15 Asp Ile Pro Pro Ala Asp Leu Ser Asp Gln Val Pro Asp Thr Glu Ser
965 970 975

20 Glu Thr Arg Ile Leu Leu Gln Gly Thr Pro Val Ala Gln Met Thr Glu
980 985 990

25 Asp Ala Val Asp Ala Glu Arg Leu Lys His Leu Ile Val Thr Pro Ser
995 1000 1005

30 Gly Cys Gly Glu Gln Asn Met Ile Gly Met Thr Pro Thr Val Ile
1010 1015 1020

Ala Val His Tyr Leu Asp Glu Thr Glu Gln Trp Glu Lys Phe Gly
1025 1030 1035

35 Leu Glu Lys Arg Gln Gly Ala Leu Glu Leu Ile Lys Lys Gly Tyr
1040 1045 1050

40 Thr Gln Gln Leu Ala Phe Arg Gln Pro Ser Ser Ala Phe Ala Ala
1055 1060 1065

45 Phe Val Lys Arg Ala Pro Ser Thr Trp Leu Thr Ala Tyr Val Val
1070 1075 1080

50 Lys Val Phe Ser Leu Ala Val Asn Leu Ile Ala Ile Asp Ser Gln
1085 1090 1095

Val Leu Cys Gly Ala Val Lys Trp Leu Ile Leu Glu Lys Gln Lys
1100 1105 1110

55 Pro Asp Gly Val Phe Gln Glu Asp Ala Pro Val Ile His Gln Glu
1115 1120 1125

60 Met Ile Gly Gly Leu Arg Asn Asn Asn Glu Lys Asp Met Ala Leu
1130 1135 1140

65 Thr Ala Phe Val Leu Ile Ser Leu Gln Glu Ala Lys Asp Ile Cys
1145 1150 1155

70 Glu Glu Gln Val Asn Ser Leu Pro Gly Ser Ile Thr Lys Ala Gly
1160 1165 1170

Asp Phe Leu Glu Ala Asn Tyr Met Asn Leu Gln Arg Ser Tyr Thr
 1175 1180 1185

5 Val Ala Ile Ala Gly Tyr Ala Leu Ala Gln Met Gly Arg Leu Lys
 1190 1195 1200

10 Gly Pro Leu Leu Asn Lys Phe Leu Thr Thr Ala Lys Asp Lys Asn
 1205 1210 1215

15 Arg Trp Glu Asp Pro Gly Lys Gln Leu Tyr Asn Val Glu Ala Thr
 1220 1225 1230

20 Ser Tyr Ala Leu Leu Ala Leu Leu Gln Leu Lys Asp Phe Asp Phe
 1235 1240 1245

25 Val Pro Pro Val Val Arg Trp Leu Asn Glu Gln Arg Tyr Tyr Gly
 1250 1255 1260

30 Gly Gly Tyr Gly Ser Thr Gln Ala Thr Phe Met Val Phe Gln Ala
 1265 1270 1275

35 Leu Ala Gln Tyr Gln Lys Asp Ala Pro Asp His Gln Glu Leu Asn
 1280 1285 1290

40 Leu Asp Val Ser Leu Gln Leu Pro Ser Arg Ser Ser Lys Ile Thr
 1295 1300 1305

45 His Arg Ile His Trp Glu Ser Ala Ser Leu Leu Arg Ser Glu Glu
 1310 1315 1320

50 Thr Lys Glu Asn Glu Gly Phe Thr Val Thr Ala Glu Gly Lys Gly
 1325 1330 1335

55 Gln Gly Thr Leu Ser Val Val Thr Met Tyr His Ala Lys Ala Lys
 1340 1345 1350

60 Asp Gln Leu Thr Cys Asn Lys Phe Asp Leu Lys Val Thr Ile Lys
 1355 1360 1365

65 Pro Ala Pro Glu Thr Glu Lys Arg Pro Gln Asp Ala Lys Asn Thr
 1370 1375 1380

70 Met Ile Leu Glu Ile Cys Thr Arg Tyr Arg Gly Asp Gln Asp Ala
 1385 1390 1395

75 Thr Met Ser Ile Leu Asp Ile Ser Met Met Thr Gly Phe Ala Pro
 1400 1405 1410

80 Asp Thr Asp Asp Leu Lys Gln Leu Ala Asn Gly Val Asp Arg Tyr
 1415 1420 1425

85 Ile Ser Lys Tyr Glu Leu Asp Lys Ala Phe Ser Asp Arg Asn Thr

	.	1430	1435	1440
5	Leu Ile Ile Tyr Leu Asp Lys Val Ser His Ser Glu Asp Asp Cys	1445	1450	1455
10	Leu Ala Phe Lys Val His Gln Tyr Phe Asn Val Glu Leu Ile Gln	1460	1465	1470
15	Pro Gly Ala Val Lys Val Tyr Ala Tyr Tyr Asn Leu Glu Glu Ser	1475	1480	1485
20	Cys Thr Arg Phe Tyr His Pro Glu Lys Glu Asp Gly Lys Leu Asn	1490	1495	1500
25	Lys Leu Cys Arg Asp Glu Leu Cys Arg Cys Ala Glu Glu Asn Cys	1505	1510	1515
30	Phe Ile Gln Lys Ser Asp Asp Lys Val Thr Leu Glu Glu Arg Leu	1520	1525	1530
35	Asp Lys Ala Cys Glu Pro Gly Val Asp Tyr Val Tyr Lys Thr Arg	1535	1540	1545
40	Leu Val Lys Val Gln Leu Ser Asn Asp Phe Asp Glu Tyr Ile Met	1550	1555	1560
45	Ala Ile Glu Gln Thr Ile Lys Ser Gly Ser Asp Glu Val Gln Val	1565	1570	1575
50	Gly Gln Gln Arg Thr Phe Ile Ser Pro Ile Lys Cys Arg Glu Ala	1580	1585	1590
55	Leu Lys Leu Glu Glu Lys Lys His Tyr Leu Met Trp Gly Leu Ser	1595	1600	1605
60	Ser Asp Phe Trp Gly Glu Lys Pro Asn Leu Ser Tyr Ile Ile Gly	1610	1615	1620
65	Lys Asp Thr Trp Val Glu His Trp Pro Glu Glu Asp Glu Cys Gln	1625	1630	1635
70	Asp Glu Glu Asn Gln Lys Gln Cys Gln Asp Leu Gly Ala Phe Thr	1640	1645	1650
60	Glu Ser Met Val Val Phe Gly Cys Pro Asn	1655	1660	
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<309> 2003-02-28
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5 <400> 45
Met Ser Gly Leu Arg Val Tyr Ser Thr Ser Val Thr Gly Ser Arg Glu
1 5 10 15

10 Ile Lys Ser Gln Gln Ser Glu Val Thr Arg Ile Leu Asp Gly Lys Arg
 20 25 30

15 Ile Gln Tyr Gln Leu Val Asp Ile Ser Gln Asp Asn Ala Leu Arg Asp
 35 40 45

20 Glu Met Arg Ala Leu Ala Gly Asn Pro Lys Ala Thr Pro Pro Gln Ile
 50 55 60

25 Val Asn Gly Asp Gln Tyr Cys Gly Asp Tyr Glu Leu Phe Val Glu Ala
 65 70 75 80

30 Val Glu Gln Asn Thr Leu Gln Glu Phe Leu Lys Leu Ala
 85 90

35 <210> 46
<211> 567
<212> PRT
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<309> 2003-02-28
<313> (1)..(567)

45 <400> 46
Met Ala Pro Leu Ala Leu His Leu Leu Val Leu Val Pro Ile Leu Leu
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50 Ser Leu Val Ala Ser Gln Asp Trp Lys Ala Glu Arg Ser Gln Asp Pro
 20 25 30

55 Phe Glu Lys Cys Met Gln Asp Pro Asp Tyr Glu Gln Leu Leu Lys Val
 35 40 45

60 Val Thr Trp Gly Leu Asn Arg Thr Leu Lys Pro Gln Arg Val Ile Val
 50 55 60

65 Val Gly Ala Gly Val Ala Gly Leu Val Ala Ala Lys Val Leu Ser Asp
 65 70 75 80

70 Ala Gly His Lys Val Thr Ile Leu Glu Ala Asp Asn Arg Ile Gly Gly
 85 90 95

75 Arg Ile Phe Thr Tyr Arg Asp Gln Asn Thr Gly Trp Ile Gly Glu Leu
 100 105 110

80 Gly Ala Met Arg Met Pro Ser Ser His Arg Ile Leu His Lys Leu Cys

115 120 125

5 Gln Gly Leu Gly Leu Asn Leu Thr Lys Phe Thr Gln Tyr Asp Lys Asn
130 135 140

10 Thr Trp Thr Glu Val His Glu Val Lys Leu Arg Asn Tyr Val Val Glu
145 150 155 160

15 Lys Val Pro Glu Lys Leu Gly Tyr Ala Leu Arg Pro Gln Glu Lys Gly
165 170 175

20 His Ser Pro Glu Asp Ile Tyr Gln Met Ala Leu Asn Gln Ala Leu Lys
180 185 190

25 Asp Leu Lys Ala Leu Gly Cys Arg Lys Ala Met Lys Lys Phe Glu Arg
195 200 205

30 His Thr Leu Leu Glu Tyr Leu Leu Gly Glu Gly Asn Leu Ser Arg Pro
210 215 220

35 Ala Val Gln Leu Leu Gly Asp Val Met Ser Glu Asp Gly Phe Phe Tyr
225 230 235 240

40 Leu Ser Phe Ala Glu Ala Leu Arg Ala His Ser Cys Leu Ser Asp Arg
245 250 255

45 Leu Gln Tyr Ser Arg Ile Val Gly Gly Trp Asp Leu Leu Pro Arg Ala
260 265 270

50 Leu Leu Ser Ser Leu Ser Gly Leu Val Leu Leu Asn Ala Pro Val Val
275 280 285

55 Ala Met Thr Gln Gly Pro His Asp Val His Val Gln Ile Glu Thr Ser
290 295 300

60 Pro Pro Ala Arg Asn Leu Lys Val Leu Lys Ala Asp Val Val Leu Leu
305 310 315 320

65 Thr Ala Ser Gly Pro Ala Val Lys Arg Ile Thr Phe Ser Pro Pro Leu
325 330 335

70 Pro Arg His Met Gln Glu Ala Leu Arg Arg Leu His Tyr Val Pro Ala
340 345 350

75 Thr Lys Val Phe Leu Ser Phe Arg Arg Pro Phe Trp Arg Glu Glu His
355 360 365

80 Ile Glu Gly Gly His Ser Asn Thr Asp Arg Pro Ser Arg Met Ile Phe
370 375 380

85 Tyr Pro Pro Pro Arg Glu Gly Ala Leu Leu Leu Ala Ser Tyr Thr Trp
385 390 395 400

Ser Asp Ala Ala Ala Ala Phe Ala Gly Leu Ser Arg Glu Glu Ala Leu
 405 410 415

5 Arg Leu Ala Leu Asp Asp Val Ala Ala Leu His Gly Pro Val Val Arg
 420 425 430

10 Gln Leu Trp Asp Gly Thr Gly Val Val Lys Arg Trp Ala Glu Asp Gln
 435 440 445

15 His Ser Gln Gly Gly Phe Val Val Gln Pro Pro Ala Leu Trp Gln Thr
 450 455 460

20 Glu Lys Asp Asp Trp Thr Val Pro Tyr Gly Arg Ile Tyr Phe Ala Gly
 465 470 475 480

25 Glu His Thr Ala Tyr Pro His Gly Trp Val Glu Thr Ala Val Lys Ser
 485 490 495

30 Ala Leu Arg Ala Ala Ile Lys Ile Asn Ser Arg Lys Gly Pro Ala Ser
 500 505 510

35 Asp Thr Ala Ser Pro Glu Gly His Ala Ser Asp Met Glu Gly Gln Gly
 515 520 525

40 His Val His Gly Val Ala Ser Ser Pro Ser His Asp Leu Ala Lys Glu
 530 535 540

45 Glu Gly Ser His Pro Pro Val Gln Gly Gln Leu Ser Leu Gln Asn Thr
 545 550 555 560

50 Thr His Thr Arg Thr Ser His
 565

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 <309> 1986-07-21
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65 <400> 47
 Met Ala Arg Val Leu Gly Ala Pro Val Ala Leu Gly Leu Trp Ser Leu
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70 Cys Trp Ser Leu Ala Ile Ala Thr Pro Leu Pro Pro Thr Ser Ala His
 20 25 30

75 Gly Asn Val Ala Glu Gly Glu Thr Lys Pro Asp Pro Asp Val Thr Glu
 35 40 45

80 Arg Cys Ser Asp Gly Trp Ser Phe Asp Ala Thr Thr Leu Asp Asp Asn

5	50	55	60	
	Gly Thr Met Leu Phe Phe Lys	Gly Glu Phe Val	Trp Lys Ser His Lys	
5	65	70	75	80
	Trp Asp Arg Glu Leu Ile Ser Glu Arg Trp Lys Asn Phe Pro Ser Pro			
10		85	90	95
	Val Asp Ala Ala Phe Arg Gln Gly His Asn Ser Val Phe Leu Ile Lys			
15	100	105	110	
	Gly Asp Lys Val Trp Val Tyr Pro Pro Glu Lys Lys Glu Lys Gly Tyr			
20	115	120	125	
	Pro Lys Leu Leu Gln Asp Glu Phe Pro Gly Ile Pro Ser Pro Leu Asp			
25	130	135	140	
	Ala Ala Val Glu Cys His Arg Gly Glu Cys Gln Ala Glu Gly Val Leu			
30	145	150	155	160
	Phe Phe Gln Gly Asp Arg Glu Trp Phe Trp Asp Leu Ala Thr Gly Thr			
35	165	170	175	
	Met Lys Glu Arg Ser Trp Pro Ala Val Gly Asn Cys Ser Ser Ala Leu			
40	180	185	190	
	Arg Trp Leu Gly Arg Tyr Tyr Cys Phe Gln Gly Asn Gln Phe Leu Arg			
45	195	200	205	
	Phe Asp Pro Val Arg Gly Glu Val Pro Pro Arg Tyr Pro Arg Asp Val			
50	210	215	220	
	Arg Asp Tyr Phe Met Pro Cys Pro Gly Arg Gly His Gly His Arg Asn			
55	225	230	235	240
	Gly Thr Gly His Gly Asn Ser Thr His His Gly Pro Glu Tyr Met Arg			
60	245	250	255	
	Cys Ser Pro His Leu Val Leu Ser Ala Leu Thr Ser Asp Asn His Gly			
65	260	265	270	
	Ala Thr Tyr Ala Phe Ser Gly Thr His Tyr Trp Arg Leu Asp Thr Ser			
70	275	280	285	
	Arg Asp Gly Trp His Ser Trp Pro Ile Ala His Gln Trp Pro Gln Gly			
75	290	295	300	
	Pro Ser Ala Val Asp Ala Ala Phe Ser Trp Glu Glu Lys Leu Tyr Leu			
80	305	310	315	320
	Val Gln Gly Thr Gln Val Tyr Val Phe Leu Thr Lys Gly Gly Tyr Thr			
85	325	330	335	

Leu Val Ser Gly Tyr Pro Lys Arg Leu Glu Lys Glu Val Gly Thr Pro
340 345 350

5 His Gly Ile Ile Leu Asp Ser Val Asp Ala Ala Phe Ile Cys Pro Gly
355 360 365

10 Ser Ser Arg Leu His Ile Met Ala Gly Arg Arg Leu Trp Trp Leu Asp
370 375 380

15 Leu Lys Ser Gly Ala Gln Ala Thr Trp Thr Glu Leu Pro Trp Pro His
385 390 395 400

20 Glu Lys Val Asp Gly Ala Leu Cys Met Glu Lys Ser Leu Gly Pro Asn
405 410 415

25 Ser Cys Ser Ala Asn Gly Pro Gly Leu Tyr Leu Ile His Gly Pro Asn
420 425 430

30 Leu Tyr Cys Tyr Ser Asp Val Glu Lys Leu Asn Ala Ala Lys Ala Leu
435 440 445

35 Pro Gln Pro Gln Asn Val Thr Ser Leu Leu Gly Cys Thr His
450 455 460

35 <210> 48
<211> 369
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<309> 1996-10-02
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45 <400> 48
Met Asp Pro Arg Lys Val Asn Glu Leu Arg Ala Phe Val Lys Met Cys
1 5 10 15

50 Lys Gln Asp Pro Ser Val Leu His Thr Glu Glu Met Arg Phe Leu Arg
20 25 30

55 Glu Trp Val Glu Ser Met Gly Gly Lys Val Pro Pro Ala Thr Gln Lys
35 40 45

60 Ala Lys Ser Glu Glu Asn Thr Lys Glu Glu Lys Pro Asp Ser Lys Lys
50 55 60

65 Val Glu Glu Asp Leu Lys Ala Asp Glu Pro Ser Ser Glu Glu Ser Asp
65 70 75 80

65 Leu Glu Ile Asp Lys Glu Gly Val Ile Glu Pro Asp Thr Asp Ala Pro
85 90 95

70 Gln Glu Met Gly Asp Glu Asn Ala Glu Ile Thr Glu Glu Met Met Asp

48

100

105

110

5 Gln Ala Asn Asp Lys Lys Val Ala Ala Ile Glu Ala Leu Asn Asp Gly
115 120 125

10 Glu Leu Gln Lys Ala Ile Asp Leu Phe Thr Asp Ala Ile Lys Leu Asn ..
130 135 140

15 Pro Arg Leu Ala Ile Leu Tyr Ala Lys Arg Ala Ser Val Phe Val Lys
145 150 155 160

20 Leu Gln Lys Pro Asn Ala Ala Ile Arg Asp Cys Asp Arg Ala Ile Glu
165 170 175

25 Ile Asn Pro Asp Ser Ala Gln Pro Tyr Lys Trp Arg Gly Lys Ala His
180 185 190

30 Arg Leu Leu Gly His Trp Glu Glu Ala Ala His Asp Leu Ala Leu Ala
195 200 205

35 Cys Lys Leu Asp Tyr Asp Glu Asp Ala Ser Ala Met Leu Lys Glu Val
210 215 220

40 Gln Pro Arg Ala Gln Lys Ile Ala Glu His Arg Arg Lys Tyr Glu Arg
225 230 235 240

45 Lys Arg Glu Glu Arg Glu Ile Lys Glu Arg Ile Glu Arg Val Lys Lys
245 250 255

50 Ala Arg Glu Glu His Glu Arg Ala Gln Arg Glu Glu Ala Arg Arg
260 265 270

55 Gln Ser Gly Ala Gln Tyr Gly Ser Phe Pro Gly Gly Phe Pro Gly Gly
275 280 285

60 Met Pro Gly Asn Phe Pro Gly Gly Met Pro Gly Met Gly Gly Met
290 295 300

65 Pro Gly Met Ala Gly Met Pro Gly Leu Asn Glu Ile Leu Ser Asp Pro
305 310 315 320

70 Glu Val Leu Ala Ala Met Gln Asp Pro Glu Val Met Val Ala Phe Gln
325 330 335

75 Asp Val Ala Gln Asn Pro Ala Asn Met Ser Lys Tyr Gln Ser Asn Pro
340 345 350

80 Lys Val Met Asn Leu Ile Ser Lys Leu Ser Ala Lys Phe Gly Gly Gln
355 360 365

85 Ala

5 <210> 49
 <211> 9
 <212> PRT
 <213> Homo sapiens

10 <400> 49

 Met Gln Leu Met His Ala Asn Ala Gln
 1 5

15 <210> 50
 <211> 9
 <212> PRT
 <213> Homo sapiens

20 <400> 50

 Leu Thr Leu Asp Ser Asn Thr Lys Tyr
 1 5

25 <210> 51
 <211> 9
 <212> PRT
 <213> Homo sapiens

30 <400> 51

 Phe Val Ile Asp Lys Ser Gly Ser Met
 1 5

35 <210> 52
 <211> 9
 <212> PRT
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40 <400> 52

 Tyr Leu Leu Asp Ser Asn Ser Trp Ile
 1 5

45 <210> 53
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 <212> PRT
 <213> Homo sapiens

50 <400> 53

 Tyr Glu Leu Asp Lys Ala Phe Ser Asp
 1 5

55 <210> 54
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60 <400> 54

 Ile Lys Ser Gln Gln Ser Glu Val Thr
 1 5

65 <210> 55
 <211> 9
 <212> PRT

<213> Homo sapiens
<400> 55
5 val Gln Ile Glu Thr Ser Pro Pro Ala
1 5

10 <210> 56
<211> 9
<212> PRT
<213> Homo sapiens

15 <400> 56
Ile Ile Leu Asp Ser Val Asp Ala Ala
1 5

20 <210> 57
<211> 9
<212> PRT
<213> Homo sapiens

25 <400> 57
Ile Glu Pro Asp Thr Asp Ala Pro Gln
1 5

30 <210> 58
<211> 17
<212> PRT
<213> Homo sapiens

35 <400> 58
Asn Ile Gln Pro Ile Phe Ala Val Thr Ser Arg Met Val Lys Thr Tyr
1 5 10 15
40
Glu

45 <210> 59
<211> 19
<212> PRT
<213> Homo sapiens

50 <400> 59
Glu Asn Asn Ile Gln Pro Ile Phe Ala Val Thr Ser Arg Met Val Lys
1 5 10 15
55
Thr Tyr Glu

60 <210> 60
<211> 17
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65 <400> 60
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Asp

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<400> 61

Tyr Pro Glu Gln Leu Lys Met Thr Val Val Lys Leu Ile Ser His Arg
1 5 10 15

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<400> 62

Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr
1 5 10 15

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<213> Homo sapiens

<400> 63

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Asn Gly Gly His Tyr Thr Tyr Ser Glu Asn Arg Val Glu Lys Asp Gly
1 5 10 15

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<211> 15
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<213> Homo sapiens

<400> 64

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Gly Pro Asn Asn Tyr Tyr Ser Phe Ala Ser Gln Gln Gln Lys Pro
1 5 10 15

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Gly Pro Asn Asn Tyr Tyr Ser Phe Ala Ser Gln Gln Gln Lys Pro Glu
1 5 10 15

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<400> 66

Gly Pro Asn Asn Tyr Tyr Ser Phe Ala Ser Gln Gln Gln Lys Pro Glu
1 5 10 15

Asp

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<400> 67

Gly Pro Asn Asn Tyr Tyr Ser Phe Ala Ser Gln Gln Gln Lys Pro Glu
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Asp Thr

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<210> 68
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<400> 68

Glu Lys Leu Trp Phe Val Pro Ala Lys Val Glu Asp Ser Gly His Tyr
1 5 10 15

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<400> 69

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Ser Leu Arg Glu Leu His Leu Asp His Asn Gln Ile Ser Arg
1 5 10

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<213> Homo sapiens

<400> 70

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Leu Arg Glu Leu His Leu Asp His Asn Gln Ile Ser Arg
1 5 10

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<210> 71
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<213> Homo sapiens

<400> 71

60

Glu Thr Met Lys Met Arg Tyr Glu His Ile Asp His Thr Phe Glu
1 5 10 15

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<210> 72
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<213> Homo sapiens

70

<400> 72

Glu Thr Met Lys Met Arg Tyr Glu His Ile Asp His Thr Phe Glu Ile
1 5 10 15

5 Gln

10 <210> 73
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15 <400> 73

His Met Phe Leu Gln Asp Glu Ile Ile Asp Lys Ser Tyr Thr Pro Ser
1 5 10 15

20 <210> 74
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<212> PRT
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25 <400> 74

Val Asp Arg Tyr Ile Ser Lys Tyr Glu Leu Asp Lys Ala Phe Ser Asp
1 5 10 15

30 Arg

35 <210> 75
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40 <400> 75

Arg Tyr Ile Ser Lys Tyr Glu Leu Asp Lys Ala Phe Ser Asp
1 5 10

45 <210> 76
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<213> Homo sapiens

50 <400> 76

55 Leu Pro Val Gly Arg Thr Val Met Val Asn Ile Glu Asn Pro Glu Gly
1 5 10 15

Ile Pro Val

60 <210> 77
<211> 20
<212> PRT
<213> Homo sapiens

65 <400> 77

70 Leu Pro Val Gly Arg Thr Val Met Val Asn Ile Glu Asn Pro Glu Gly
1 5 10 15

Ile Pro Val Lys
20

5

<210> 78
<211> 18
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10 <213> Homo sapiens
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Gly Thr Pro His Gly Ile Ile Leu Asp Ser Val Asp Ala Ala Phe Ile
15 1 5 10 15

Cys Pro

20

<210> 79
<211> 18
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25 <213> Homo sapiens
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Met Ala Thr Pro Leu Leu Met Gln Ala Leu Pro Met Gly Ala Leu Pro
30 1 5 10 15

Gln Gly

35

<210> 80
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40 <213> Homo sapiens
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Met Ala Thr Pro Leu Leu Met Gln Ala Leu Pro Met Gly Ala Leu Pro
45 1 5 10 15

Gln

50

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55 <213> Homo sapiens
<400> 81

Met Ala Thr Pro Leu Leu Met Gln Ala Leu Pro Met Gly Ala Leu Pro
60 1 5 10 15

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65 <213> Homo sapiens
<400> 82

70 Met Ala Thr Pro Leu Leu Met Gln Ala Leu Pro Met Gly Ala Leu Pro

1 5 10 15

5 Gln Gly Pro Met Gln
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10 <210> 83
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15 <400> 83
Ala Thr Pro Leu Leu Met Gln Ala Leu Pro Met Gly Ala Leu Pro
1 5 10 15

20 <210> 84
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25 <400> 84
His Pro Pro Val Gln Trp Ala Phe Gln Glu Thr Ser Val Glu Ser Ala
1 5 10 15

30 Val Asp Thr Pro Phe Pro Ala
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40 <400> 85
His Pro Pro Val Gln Trp Ala Phe Gln Glu Thr Ser Val Glu Ser Ala
1 5 10 15

45 Val Asp Thr Pro Phe Pro Ala Gly
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His Pro Pro Val Gln Trp Ala Phe Gln Glu Thr Ser Val Glu Ser Ala
1 5 10 15

60 Val Asp Thr Pro Phe Pro
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Lys Ile Tyr Leu Tyr Thr Leu Asn Asp Asn Ala Arg Ser Ser Pro Val
1 5 10 15

5 Val

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15 <400> 88

Lys Ile Tyr Leu Tyr Thr Leu Asn Asp Asn Ala Arg Ser Ser Pro Val
1 5 10 15

20 <210> 89
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25 <400> 89

Ile Tyr Leu Tyr Thr Leu Asn Asp Asn Ala Arg Ser Ser Pro Val Val
1 5 10 15

30 Ile

35 <210> 90
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40 <400> 90

Ile Tyr Leu Tyr Thr Leu Asn Asp Asn Ala Arg Ser Ser Pro Val
1 5 10 15

45 <210> 91
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50 <400> 91

Ile Tyr Leu Tyr Thr Leu Asn Asp Asn Ala Arg Ser Ser Pro Val Val
1 5 10 15

55 <210> 92
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60 <400> 92

Tyr Asn Ser Tyr Ser Val Ser Asn Ser Glu Lys Asp Ile Met Ala
1 5 10 15

65 <210> 93
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<212> PRT

<213> Homo sapiens
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5 Ala Gly Ser Leu Thr Leu Ser Lys Thr Glu Leu Gly Lys Lys Ala
1 5 10 15

10 <210> 94
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<213> Homo sapiens
<400> 94

15 Ala Gly Ser Leu Thr Leu Ser Lys Thr Glu Leu Gly Lys Lys Ala Asp
1 5 10 15

20 <210> 95
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<213> Homo sapiens
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25 Val Pro Lys Asp Tyr Thr Gly Glu Asp Val Thr Pro Gln Asn
1 5 10

30 <210> 96
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35 Asp Ser Lys Phe His Gln Ala Ile Asn Asp Ala His Gln
1 5 10

40 <210> 97
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45 Met Pro Leu Glu Phe Lys Thr Leu Asn Val Leu His Asn Arg Gly
1 5 10 15

50 <210> 98
<211> 15
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<400> 98

55 Ala Thr Arg Ser Ile Gln Val Asp Gly Lys Thr Ile Lys Ala Gln
1 5 10 15

60 <210> 99
<211> 18
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65 70

Ala Thr Arg Ser Ile Gln Val Asp Gly Lys Thr Ile Lys Ala Gln Ile
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5 Trp Asp

10 <210> 100
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<212> PRT
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15 <400> 100
Thr Arg Ser Ile Gln Val Asp Gly Lys Thr Ile Lys Ala Gln
1 5 10

20 <210> 101
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25 <400> 101
Ala Thr Arg Ser Ile Gln Val Asp Gly Lys Thr Ile Lys Ala Gln Ile
1 5 10 15

30 Trp

35 <210> 102
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40 <400> 102
Arg Ser Ile Gln Val Asp Gly Lys Thr Ile Lys Ala Gln
1 5 10

45 <210> 103
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50 <400> 103
Ile Gln Pro Ile Phe Ala Val Thr Ser
1 5

55 <210> 104
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60 <400> 104
Val Phe Gly Glu Asp Ser Val Gly Val
1 5

65 <210> 105
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5 Leu Lys Met Thr Val Val Lys Leu Ile
1 5

10 <210> 106
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<400> 106
15 Tyr Leu Gln Met Asn Ser Leu Arg Ala
1 5

20 <210> 107
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<400> 107
Tyr Thr Tyr Ser Glu Asn Arg Val Glu
1 5

30 <210> 108
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35 <400> 108
Tyr Tyr Ser Phe Ala Ser Gln Gln Gln
1 5
40

45 <210> 109
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Phe Val Pro Ala Lys Val Glu Asp Ser
1 5

55 <210> 110
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<400> 110
60 Leu His Leu Asp His Asn Gln Ile Ser
1 5

65 <210> 111
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70

Met Arg Tyr Glu His Ile Asp His Thr
1 5

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10 <400> 112

Phe Leu Gln Asp Glu Ile Ile Asp Lys
1 5

15 <210> 113
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<212> PRT
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20 <400> 113

Met Val Asn Ile Glu Asn Pro Glu Gly
1 5

25 <210> 114
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30 <400> 114

Leu Met Gln Ala Leu Pro Met Gly Ala
1 5

35 <210> 115
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<212> PRT
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40 <400> 115

Trp Ala Phe Gln Glu Thr Ser Val Glu
1 5

45 <210> 116
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<212> PRT
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50 <400> 116

Tyr Thr Leu Asn Asp Asn Ala Arg Ser

1 5

55 <210> 117
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60 <400> 117

Tyr Ser Val Ser Asn Ser Glu Lys Asp
1 5

5 <210> 118
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 <212> PRT
 <213> Homo sapiens

10 <400> 118

 Leu Thr Leu Ser Lys Thr Glu Leu Gly
 1 5

15 <210> 119
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 <400> 119

20 Tyr Thr Gly Glu Asp Val Thr Pro Gln
 1 5

25 <210> 120
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 <212> PRT
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 <400> 120

30 Phe His Gln Ala Ile Asn Asp Ala His
 1 5

35 <210> 121
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 <400> 121

40 Phe Lys Thr Leu Asn Val Leu His Asn
 1 5

45 <210> 122
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 Ile Gln Val Asp Gly Lys Thr Ile Lys
 1 5

55 <210> 123
 <211> 769
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 Met Leu Gly Leu Arg Pro Pro Leu Leu Ala Leu Val Gly Leu Leu Ser
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Leu Gly Cys Val Leu Ser Gln Glu Cys Thr Lys Phe Lys Val Ser Ser
 20 25 30

5 Cys Arg Glu Cys Ile Glu Ser Gly Pro Gly Cys Thr Trp Cys Gln Lys
 35 40 45

10 Leu Asn Phe Thr Gly Pro Gly Asp Pro Asp Ser Ile Arg Cys Asp Thr
 50 55 60

15 Arg Pro Gln Leu Leu Met Arg Gly Cys Ala Ala Asp Asp Ile Met Asp
 65 70 75 80

20 Pro Thr Ser Leu Ala Glu Thr Gln Glu Asp His Asn Gly Gly Gln Lys
 85 90 95

25 Gln Leu Ser Pro Gln Lys Val Thr Leu Tyr Leu Arg Pro Gly Gln Ala
 100 105 110

30 Ala Ala Phe Asn Val Thr Phe Arg Arg Ala Lys Gly Tyr Pro Ile Asp
 115 120 125

35 Leu Tyr Tyr Leu Met Asp Leu Ser Tyr Ser Met Leu Asp Asp Leu Arg
 130 135 140

40 Asn Val Lys Lys Leu Gly Gly Asp Leu Leu Arg Ala Leu Asn Glu Ile
 145 150 155 160

45 Thr Glu Ser Gly Arg Ile Gly Phe Gly Ser Phe Val Asp Lys Thr Val
 165 170 175

50 Leu Pro Phe Val Asn Thr His Pro Asp Lys Leu Arg Asn Pro Cys Pro
 180 185 190

55 Asn Lys Glu Lys Glu Cys Gln Pro Pro Phe Ala Phe Arg His Val Leu
 195 200 205

60 Lys Leu Thr Asn Asn Ser Asn Gln Phe Gln Thr Glu Val Gly Lys Gln
 210 215 220

65 Leu Ile Ser Gly Asn Leu Asp Ala Pro Glu Gly Gly Leu Asp Ala Met
 225 230 235 240

70 Met Gln Val Ala Ala Cys Pro Glu Glu Ile Gly Trp Arg Asn Val Thr
 245 250 255

75 Arg Leu Leu Val Phe Ala Thr Asp Asp Gly Phe His Phe Ala Gly Asp
 260 265 270

80 Gly Lys Leu Gly Ala Ile Leu Thr Pro Asn Asp Gly Arg Cys His Leu
 275 280 285

85 Glu Asp Asn Leu Tyr Lys Arg Ser Asn Glu Phe Asp Tyr Pro Ser Val

290 295 300

5 Gly Gln Leu Ala His Lys Leu Ala Glu Asn Asn Ile Gln Pro Ile Phe
305 310 315 320

10 Ala Val Thr Ser Arg Met Val Lys Thr Tyr Glu Lys Leu Thr Glu Ile
325 330 335

15 Ile Pro Lys Ser Ala Val Gly Glu Leu Ser Glu Asp Ser Ser Asn Val
340 345 350

20 Val His Leu Ile Lys Asn Ala Tyr Asn Lys Leu Ser Ser Arg Val Phe
355 360 365

25 Leu Asp His Asn Ala Leu Pro Asp Thr Leu Lys Val Thr Tyr Asp Ser
370 375 380

30 Phe Cys Ser Asn Gly Val Thr His Arg Asn Gln Pro Arg Gly Asp Cys
385 390 395 400

35 Asp Gly Val Gln Ile Asn Val Pro Ile Thr Phe Gln Val Lys Val Thr
405 410 415

40 Ala Thr Glu Cys Ile Gln Glu Gln Ser Phe Val Ile Arg Ala Leu Gly
420 425 430

45 Phe Thr Asp Ile Val Thr Val Gln Val Leu Pro Gln Cys Glu Cys Arg
435 440 445

50 Cys Arg Asp Gln Ser Arg Asp Arg Ser Leu Cys His Gly Lys Gly Phe
450 455 460

55 Leu Glu Cys Gly Ile Cys Arg Cys Asp Thr Gly Tyr Ile Gly Lys Asn
465 470 475 480

60 Cys Glu Cys Gln Thr Gln Gly Arg Ser Ser Gln Glu Leu Glu Gly Ser
485 490 495

65 Cys Arg Lys Asp Asn Asn Ser Ile Ile Cys Ser Gly Leu Gly Asp Cys
500 505 510

70 Val Cys Gly Gln Cys Leu Cys His Thr Ser Asp Val Pro Gly Lys Leu
515 520 525

75 Ile Tyr Gly Gln Tyr Cys Glu Cys Asp Thr Ile Asn Cys Glu Arg Tyr
530 535 540

80 Asn Gly Gln Val Cys Gly Gly Pro Gly Arg Gly Leu Cys Phe Cys Gly
545 550 555 560

85 Lys Cys Arg Cys His Pro Gly Phe Glu Gly Ser Ala Cys Gln Cys Glu
565 570 575

Arg Thr Thr Glu Gly Cys Leu Asn Pro Arg Arg Val Glu Cys Ser Gly
 580 585 590

5 Arg Gly Arg Cys Arg Cys Asn Val Cys Glu Cys His Ser Gly Tyr Gln
 595 600 605

10 Leu Pro Leu Cys Gln Glu Cys Pro Gly Cys Pro Ser Pro Cys Gly Lys
 610 615 620

15 Tyr Ile Ser Cys Ala Glu Cys Leu Lys Phe Glu Lys Gly Pro Phe Gly
 625 630 635 640

Lys Asn Cys Ser Ala Ala Cys Pro Gly Leu Gln Leu Ser Asn Asn Pro
 645 650 655

20 Val Lys Gly Arg Thr Cys Lys Glu Arg Asp Ser Glu Gly Cys Trp Val
 660 665 670

25 Ala Tyr Thr Leu Glu Gln Gln Asp Gly Met Asp Arg Tyr Leu Ile Tyr
 675 680 685

30 Val Asp Glu Ser Arg Glu Cys Val Ala Gly Pro Asn Ile Ala Ala Ile
 690 695 700

35 Val Gly Gly Thr Val Ala Gly Ile Val Leu Ile Gly Ile Leu Leu Leu
 705 710 715 720

Val Ile Trp Lys Ala Leu Ile His Leu Ser Asp Leu Arg Glu Tyr Arg
 725 730 735

40 Arg Phe Glu Lys Glu Lys Leu Lys Ser Gln Trp Asn Asn Asp Asn Pro
 740 745 750

45 Leu Phe Lys Ser Ala Thr Thr Thr Val Met Asn Pro Lys Phe Ala Glu
 755 760 765

50 Ser

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Met Cys Phe Ser Phe Ile Met Pro Pro Ala Met Ala Asp Ile Leu Asp
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70 Ile Trp Ala Val Asp Ser Gln Ile Ala Ser Asp Gly Ser Ile Pro Val

65

20

25

30

Asp Phe Leu Leu Pro Thr Gly Ile Tyr Ile Gln Leu Glu Val Pro Arg
5 35 40 45

Glu Ala Thr Ile Ser Tyr Ile Lys Gln Met Leu Trp Lys Gln Val His
10 50 55 60

Asn Tyr Pro Met Phe Asn Leu Leu Met Asp Ile Asp Ser Tyr Met Phe
15 65 70 75 80

Ala Cys Val Asn Gln Thr Ala Val Tyr Glu Glu Leu Glu Asp Glu Thr
85 90 95

Arg Arg Leu Cys Asp Val Arg Pro Phe Leu Pro Val Leu Lys Leu Val
20 100 105 110

Thr Arg Ser Cys Asp Pro Gly Glu Lys Leu Asp Ser Lys Ile Gly Val
25 115 120 125

Leu Ile Gly Lys Gly Leu His Glu Phe Asp Ser Leu Lys Asp Pro Glu
30 130 135 140

Val Asn Glu Phe Arg Arg Lys Met Arg Lys Phe Ser Glu Glu Lys Ile
35 145 150 155 160

Leu Ser Leu Val Gly Leu Ser Trp Met Asp Trp Leu Lys Gln Thr Tyr
40 165 170 175

Pro Pro Glu His Glu Pro Ser Ile Pro Glu Asn Leu Glu Asp Lys Leu
45 180 185 190

Tyr Gly Gly Lys Leu Ile Val Ala Val His Phe Glu Asn Cys Gln Asp
50 195 200 205

Val Phe Ser Phe Gln Val Ser Pro Asn Met Asn Pro Ile Lys Val Asn
55 210 215 220

Glu Leu Ala Ile Gln Lys Arg Leu Thr Ile His Gly Lys Glu Asp Glu
60 225 230 235 240

Val Ser Pro Tyr Asp Tyr Val Leu Gln Val Ser Gly Arg Val Glu Tyr
65 245 250 255

Val Phe Gly Asp His Pro Leu Ile Gln Phe Gln Tyr Ile Arg Asn Cys
70 260 265 270

Val Met Asn Arg Ala Leu Pro His Phe Ile Leu Val Glu Cys Cys Lys
75 275 280 285

Ile Lys Lys Met Tyr Glu Gln Glu Met Ile Ala Ile Glu Ala Ala Ile
80 290 295 300

Asn Arg Asn Ser Ser Asn Leu Pro Leu Pro Leu Pro Pro Lys Lys Thr
305 310 315 320

5 Arg Ile Ile Ser His Val Trp Glu Asn Asn Asn Pro Phe Gln Ile Val
325 330 335

10 Leu Val Lys Gly Asn Lys Leu Asn Thr Glu Glu Thr Val Lys Val His
340 345 350

15 Val Arg Ala Gly Leu Phe His Gly Thr Glu Leu Leu Cys Lys Thr Ile
355 360 365

20 Val Ser Ser Glu Val Ser Gly Lys Asn Asp His Ile Trp Asn Glu Pro
370 375 380

25 Leu Glu Phe Asp Ile Asn Ile Cys Asp Leu Pro Arg Met Ala Arg Leu
385 390 395 400

30 Cys Phe Ala Val Tyr Ala Val Leu Asp Lys Val Lys Thr Lys Lys Ser
405 410 415

35 Thr Lys Thr Ile Asn Pro Ser Lys Tyr Gln Thr Ile Arg Lys Ala Gly
420 425 430

40 Lys Val His Tyr Pro Val Ala Trp Val Asn Thr Met Val Phe Asp Phe
435 440 445

45 Lys Gly Gln Leu Arg Thr Gly Asp Ile Ile Leu His Ser Trp Ser Ser
450 455 460

50 Phe Pro Asp Glu Leu Glu Glu Met Leu Asn Pro Met Gly Thr Val Gln
465 470 475 480

55 Thr Asn Pro Tyr Thr Glu Asn Ala Thr Ala Leu His Val Lys Phe Pro
485 490 495

60 Glu Asn Lys Lys Gln Pro Tyr Tyr Tyr Pro Pro Phe Asp Lys Ile Ile
500 505 510

65 Glu Lys Ala Ala Glu Ile Ala Ser Ser Asp Ser Ala Asn Val Ser Ser
515 520 525

70 Arg Gly Gly Lys Lys Phe Leu Pro Val Leu Lys Glu Ile Leu Asp Arg
530 535 540

Asp Pro Leu Ser Gln Leu Cys Glu Asn Glu Met Asp Leu Ile Trp Thr
545 550 555 560

Leu Arg Gln Asp Cys Arg Glu Ile Phe Pro Gln Ser Leu Pro Lys Leu
565 570 575

70 Leu Leu Ser Ile Lys Trp Asn Lys Leu Glu Asp Val Ala Gln Leu Gln

	580	585	590	
5	Ala Leu Leu Gln Ile Trp Pro Lys	Leu Pro Pro Arg Glu	Ala Leu Glu	
	595	600	605	
10	Leu Leu Asp Phe Asn Tyr Pro Asp Gln Tyr Val Arg Glu Tyr Ala Val			
	610	615	620	
15	Gly Cys Leu Arg Gln Met Ser Asp Glu Glu	Leu Ser Gln Tyr Leu	Leu	
	625	630	635	640
20	Gln Leu Val Gln Val Leu Lys Tyr Glu Pro Phe Leu Asp Cys Ala Leu			
	645	650	655	
25	Ser Arg Phe Leu Leu Glu Arg Ala Leu Gly Asn Arg Arg Ile Gly Gln			
	660	665	670	
30	Phe Leu Phe Trp His Leu Arg Ser Glu Val His Ile Pro Ala Val Ser			
	675	680	685	
35	Val Gln Phe Gly Val Ile Leu Glu Ala Tyr Cys Arg Gly Ser Val Gly			
	690	695	700	
40	His Met Lys Val Leu Ser Lys Gln Val Glu Ala Leu Asn Lys Leu Lys			
	705	710	715	720
45	Thr Leu Asn Ser Leu Ile Lys Leu Asn Ala Val Lys Leu Asn Arg Ala			
	725	730	735	
50	Lys Gly Lys Glu Ala Met His Thr Cys Leu Lys Gln Ser Ala Tyr Arg			
	740	745	750	
55	Glu Ala Leu Ser Asp Leu Gln Ser Pro Leu Asn Pro Cys Val Ile Leu			
	755	760	765	
60	Ser Glu Leu Tyr Val Glu Lys Cys Lys Tyr Met Asp Ser Lys Met Lys			
	770	775	780	
65	Pro Leu Trp Leu Val Tyr Asn Asn Lys Val Phe Gly Glu Asp Ser Val			
	785	790	795	800
70	Gly Val Ile Phe Lys Asn Gly Asp Asp Leu Arg Gln Asp Met Leu Thr			
	805	810	815	
75	Leu Gln Met Leu Arg Leu Met Asp Leu Leu Trp Lys Glu Ala Gly Leu			
	820	825	830	
80	Asp Leu Arg Met Leu Pro Tyr Gly Cys Leu Ala Thr Gly Asp Arg Ser			
	835	840	845	
85	Gly Leu Ile Glu Val Val Ser Thr Ser Glu Thr Ile Ala Asp Ile Gln			
	850	855	860	

Leu Asn Ser Ser Asn Val Ala Ala Ala Ala Ala Phe Asn Lys Asp Ala
 865 870 875 880

5 Leu Leu Asn Trp Leu Lys Glu Tyr Asn Ser Gly Asp Asp Leu Asp Arg
 885 890 895

10 Ala Ile Glu Glu Phe Thr Leu Ser Cys Ala Gly Tyr Cys Val Ala Ser
 900 905 910

15 Tyr Val Leu Gly Ile Gly Asp Arg His Ser Asp Asn Ile Met Val Lys
 915 920 925

20 Lys Thr Gly Gln Leu Phe His Ile Asp Phe Gly His Ile Leu Gly Asn
 930 935 940

25 Phe Lys Ser Lys Phe Gly Ile Lys Arg Glu Arg Val Pro Phe Ile Leu
 945 950 955 960

30 Thr Tyr Asp Phe Ile His Val Ile Gln Gln Gly Lys Thr Gly Asn Thr
 965 970 975

35 Glu Lys Phe Gly Arg Phe Arg Gln Cys Cys Glu Asp Ala Tyr Leu Ile
 980 985 990

40 Leu Arg Arg His Gly Asn Leu Phe Ile Thr Leu Phe Ala Leu Met Leu
 995 1000 1005

45 Thr Ala Gly Leu Pro Glu Leu Thr Ser Val Lys Asp Ile Gln Tyr
 1010 1015 1020

50 Leu Lys Asp Ser Leu Ala Leu Gly Lys Ser Glu Glu Ala Leu
 1025 1030 1035

55 Lys Gln Phe Lys Gln Lys Phe Asp Glu Ala Leu Arg Glu Ser Trp
 1040 1045 1050

60 Thr Thr Lys Val Asn Trp Met Ala His Thr Val Arg Lys Asp Tyr
 1055 1060 1065

65 Arg Ser
 1070

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 <313> (1)..(431)

70 <400> 125
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Asp Ser Lys Gly Ser Asn Glu Leu His Gln Val Pro Ser Asn Cys Asp
5 20 25 30

Cys Leu Asn Gly Gly Thr Cys Val Ser Asn Lys Tyr Phe Ser Asn Ile
10 35 40 45

His Trp Cys Asn Cys Pro Lys Lys Phe Gly Gly Gln His Cys Glu Ile
15 50 55 60

Asp Lys Ser Lys Thr Cys Tyr Glu Gly Asn Gly His Phe Tyr Arg Gly
20 65 70 75 80

Lys Ala Ser Thr Asp Thr Met Gly Arg Pro Cys Leu Pro Trp Asn Ser
25 85 90 95

Ala Thr Val Leu Gln Gln Thr Tyr His Ala His Arg Ser Asp Ala Leu
30 100 105 110

Gln Leu Gly Leu Gly Lys His Asn Tyr Cys Arg Asn Pro Asp Asn Arg
35 115 120 125

Arg Arg Pro Trp Cys Tyr Val Gln Val Gly Leu Lys Pro Leu Val Gln
40 130 135 140

Glu Cys Met Val His Asp Cys Ala Asp Gly Lys Lys Pro Ser Ser Pro
45 145 150 155 160

Pro Glu Glu Leu Lys Phe Gln Cys Gly Gln Lys Thr Leu Arg Pro Arg
50 165 170 175

Phe Lys Ile Ile Gly Gly Glu Phe Thr Thr Ile Glu Asn Gln Pro Trp
55 180 185 190

Phe Ala Ala Ile Tyr Arg Arg His Arg Gly Gly Ser Val Thr Tyr Val
60 195 200 205

Cys Gly Gly Ser Leu Met Ser Pro Cys Trp Val Ile Ser Ala Thr His
65 210 215 220

Cys Phe Ile Asp Tyr Pro Lys Lys Glu Asp Tyr Ile Val Tyr Leu Gly
70 225 230 235 240

Arg Ser Arg Leu Asn Ser Asn Thr Gln Gly Glu Met Lys Phe Glu Val.
75 245 250 255

Glu Asn Leu Ile Leu His Lys Asp Tyr Ser Ala Asp Thr Leu Ala His
80 260 265 270

His Asn Asp Ile Ala Leu Leu Lys Ile Arg Ser Lys Glu Gly Arg Cys
85 275 280 285

Ala Gln Pro Ser Arg Thr Ile Gln Thr Ile Cys Leu Pro Ser Met Tyr
 290 295 300

5 Asn Asp Pro Gln Phe Gly Thr Ser Cys Glu Ile Thr Gly Phe Gly Lys
 305 310 315 320

10 Glu Asn Ser Thr Asp Tyr Leu Tyr Pro Glu Gln Leu Lys Met Thr Val
 325 330 335

15 Val Lys Leu Ile Ser His Arg Glu Cys Gln Gln Pro His Tyr Tyr Gly
 340 345 350

20 Ser Glu Val Thr Thr Lys Met Leu Cys Ala Ala Asp Pro Gln Trp Lys
 355 360 365

25 Thr Asp Ser Cys Gln Gly Asp Ser Gly Gly Pro Leu Val Cys Ser Leu
 370 375 380

30 Gln Gly Arg Met Thr Leu Thr Gly Ile Val Ser Trp Gly Arg Gly Cys
 385 390 395 400

35 Ala Leu Lys Asp Lys Pro Gly Val Tyr Thr Arg Val Ser His Phe Leu
 405 410 415

40 Pro Trp Ile Arg Ser His Thr Lys Glu Glu Asn Gly Leu Ala Leu
 420 425 430

<210> 126
 <211> 117
 <212> PRT
 <213> Homo sapiens

45 <300>
 <308> Swissprot/P01764
 <309> 1986-07-21
 <313> (1)..(117)

50 <400> 126
 Met Glu Phe Gly Leu Ser Trp Leu Phe Leu Val Ala Ile Leu Lys Gly
 1 5 10 15

55 Val Gln Cys Glu Val Gln Leu Leu Glu Ser Gly Gly Leu Val Gln
 20 25 30

60 Pro Gly Gly Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe
 35 40 45

65 Ser Ser Tyr Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu
 50 55 60

70 Glu Trp Val Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Gly
 65 70 75 80

75 Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn

85 90 95
Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val
105 110

Lys

iens

t/Q99497
01
9)

Arg Ala Leu Val Ile Leu Ala Lys Gly Ala Glu Glu
5 10 15

Ile Pro Val Asp Val Met Arg Arg Ala Gly Ile Lys
25 30

Gly Leu Ala Gly Lys Asp Pro Val Gln Cys Ser Arg
40 45

Cys Pro Asp Ala Ser Leu Glu Asp Ala Lys Lys Glu
55 60

Val Val Val Leu Pro Gly Gly Asn Leu Gly Ala Gln
70 75 80

Ser Ala Ala Val Lys Glu Ile Leu Lys Glu Gln Glu
85 90 95

Leu Ile Ala Ala Ile Cys Ala Gly Pro Thr Ala Leu
105 110

Ile Gly Phe Gly Ser Lys Val Thr Thr His Pro Leu
120 125

Met Met Asn Gly Gly His Tyr Thr Tyr Ser Glu Asn
135 140

Asp Gly Leu Ile Leu Thr Ser Arg Gly Pro Gly Thr
150 155 160

Ala Leu Ala Ile Val Glu Ala Leu Asn Gly Lys Glu
165 170 175

Val Lys Ala Pro Leu Val Leu Lys Asp
185

5 <210> 128
 <211> 257
 <212> PRT
 <213> Homo sapiens

10 <300>
 <308> Swissprot/P48556
 <309> 1996-02-01
 <313> (1)..(257)

15 <400> 128

Met Tyr Glu Gln Leu Lys Gly Glu Trp Asn Arg Lys Ser Pro Asn Leu
1 5 10 15

20 Ser Lys Cys Gly Glu Glu Leu Gly Arg Leu Lys Leu Val Leu Leu Glu
 20 25 30

25 Leu Asn Phe Leu Pro Thr Thr Gly Thr Lys Leu Thr Lys Gln Gln Leu
 35 40 45

30 Ile Leu Ala Arg Asp Ile Leu Glu Ile Gly Ala Gln Trp Ser Ile Leu
 50 55 60

35 Arg Lys Asp Ile Pro Ser Phe Glu Arg Tyr Met Ala Gln Leu Lys Cys
 65 70 75 80

40 Tyr Tyr Phe Asp Tyr Lys Glu Gln Leu Pro Glu Ser Ala Tyr Met His
 85 90 95

45 Gln Leu Leu Gly Leu Asn Leu Leu Phe Leu Leu Ser Gln Asn Arg Val
 100 105 110

50 Ala Glu Phe His Thr Glu Leu Glu Arg Leu Pro Ala Lys Asp Ile Gln
 115 120 125

55 Thr Asn Val Tyr Ile Lys His Pro Val Ser Leu Glu Gln Tyr Leu Met
 130 135 140

60 Glu Gly Ser Tyr Asn Lys Val Phe Leu Ala Lys Gly Asn Ile Pro Ala
 145 150 155 160

65 Glu Ser Tyr Thr Phe Phe Ile Asp Ile Leu Leu Asp Thr Ile Arg Asp
 165 170 175

70 Glu Ile Ala Gly Cys Ile Glu Lys Ala Tyr Glu Lys Ile Leu Phe Thr
 180 185 190

75 Glu Ala Thr Arg Ile Leu Phe Phe Asn Thr Pro Lys Lys Met Thr Asp
 195 200 205

80 Tyr Ala Lys Lys Arg Gly Trp Val Leu Gly Pro Asn Asn Tyr Tyr Ser
 210 215 220

85 Phe Ala Ser Gln Gln Gln Lys Pro Glu Asp Thr Thr Ile Pro Ser Thr

225 230 235 240

5 Glu Leu Ala Lys Gln Val Ile Glu Tyr Ala Arg Gln Leu Glu Met Ile
245 250 255

val

10

<210> 129
<211> 569

15 <212> PRT
<213> Homo sapiens

<300>
<308> Swissprot/P14778
<309> 1990-04-01
20 <313> (1)..(569)

<400> 129

25 Met Lys Val Leu Leu Arg Leu Ile Cys Phe Ile Ala Leu Leu Ile Ser
1 5 10 15

Ser Leu Glu Ala Asp Lys Cys Lys Glu Arg Glu Glu Lys Ile Ile Leu
20 25 30

30 Val Ser Ser Ala Asn Glu Ile Asp Val Arg Pro Cys Pro Leu Asn Pro
35 40 45

35 Asn Glu His Lys Gly Thr Ile Thr Trp Tyr Lys Asp Asp Ser Lys Thr
50 55 60

40 Pro Val Ser Thr Glu Gln Ala Ser Arg Ile His Gln His Lys Glu Lys
65 70 75 80

45 Leu Trp Phe Val Pro Ala Lys Val Glu Asp Ser Gly His Tyr Tyr Cys
85 90 95

50 Val Val Arg Asn Ser Ser Tyr Cys Leu Arg Ile Lys Ile Ser Ala Lys
100 105 110

55 Phe Val Glu Asn Glu Pro Asn Leu Cys Tyr Asn Ala Gln Ala Ile Phe
115 120 125

Lys Gln Lys Leu Pro Val Ala Gly Asp Gly Gly Leu Val Cys Pro Tyr
130 135 140

60 Met Glu Phe Phe Lys Asn Glu Asn Asn Glu Leu Pro Lys Leu Gln Trp
145 150 155 160

65 Tyr Lys Asp Cys Lys Pro Leu Leu Leu Asp Asn Ile His Phe Ser Gly
165 170 175

70 Val Lys Asp Arg Leu Ile Val Met Asn Val Ala Glu Lys His Arg Gly
180 185 190

Asn Tyr Thr Cys His Ala Ser Tyr Thr Tyr Leu Gly Lys Gln Tyr Pro
195 200 205

5 Ile Thr Arg Val Ile Glu Phe Ile Thr Leu Glu Glu Asn Lys Pro Thr
210 215 220

10 Arg Pro Val Ile Val Ser Pro Ala Asn Glu Thr Met Glu Val Asp Leu
225 230 235 240

15 Gly Ser Gln Ile Gln Leu Ile Cys Asn Val Thr Gly Gln Leu Ser Asp
245 250 255

Ile Ala Tyr Trp Lys Trp Asn Gly Ser Val Ile Asp Glu Asp Asp Pro
260 265 270

20 Val Leu Gly Glu Asp Tyr Tyr Ser Val Glu Asn Pro Ala Asn Lys Arg
275 280 285

25 Arg Ser Thr Leu Ile Thr Val Leu Asn Ile Ser Glu Ile Glu Ser Arg
290 295 300

30 Phe Tyr Lys His Pro Phe Thr Cys Phe Ala Lys Asn Thr His Gly Ile
305 310 315 320

35 Asp Ala Ala Tyr Ile Gln Leu Ile Tyr Pro Val Thr Asn Phe Gln Lys
325 330 335

His Met Ile Gly Ile Cys Val Thr Leu Thr Val Ile Ile Val Cys Ser
340 345 350

40 Val Phe Ile Tyr Lys Ile Phe Lys Ile Asp Ile Val Leu Trp Tyr Arg
355 360 365

45 Asp Ser Cys Tyr Asp Phe Leu Pro Ile Lys Ala Ser Asp Gly Lys Thr
370 375 380

50 Tyr Asp Ala Tyr Ile Leu Tyr Pro Lys Thr Val Gly Glu Gly Ser Thr
385 390 395 400

55 Ser Asp Cys Asp Ile Phe Val Phe Lys Val Leu Pro Glu Val Leu Glu
405 410 415

Lys Gln Cys Gly Tyr Lys Leu Phe Ile Tyr Gly Arg Asp Asp Tyr Val
420 425 430

60 Gly Glu Asp Ile Val Glu Val Ile Asn Glu Asn Val Lys Lys Ser Arg
435 440 445

65 Arg Leu Ile Ile Ile Leu Val Arg Glu Thr Ser Gly Phe Ser Trp Leu
450 455 460

70 Gly Gly Ser Ser Glu Glu Gln Ile Ala Met Tyr Asn Ala Leu Val Gln

75

465

470

475

480

Asp Gly Ile Lys Val Val Leu Leu Glu Leu Glu Lys Ile Gln Asp Tyr
 5 485 490 495

Glu Lys Met Pro Glu Ser Ile Lys Phe Ile Lys Gln Lys His Gly Ala
 10 500 505 510

Ile Arg Trp Ser Gly Asp Phe Thr Gln Gly Pro Gln Ser Ala Lys Thr
 515 520 525

Arg Phe Trp Lys Asn Val Arg Tyr His Met Pro Val Gln Arg Arg Ser
 15 530 535 540

Pro Ser Ser Lys His Gln Leu Leu Ser Pro Ala Thr Lys Glu Lys Leu
 20 545 550 555 560

Gln Arg Glu Ala His Val Pro Leu Gly
 25 565

<210> 130

<211> 376

<212> PRT

<213> Homo sapiens

<300>

<308> Swissprot/Q06828

<309> 1994-06-01

<313> (1)..(376)

<400> 130

Met Gln Trp Thr Ser Leu Leu Leu Leu Ala Gly Leu Phe Ser Leu Ser
 40 1 5 10 15

Gln Ala Gln Tyr Glu Asp Asp Pro His Trp Trp Phe His Tyr Leu Arg
 45 20 25 30

Ser Gln Gln Ser Thr Tyr Tyr Asp Pro Tyr Asp Pro Tyr Pro Tyr Glu
 50 35 40 45

Thr Tyr Glu Pro Tyr Pro Tyr Gly Val Asp Glu Gly Pro Ala Tyr Thr
 55 50 55 60

Tyr Gly Ser Pro Ser Pro Pro Asp Pro Arg Asp Cys Pro Gln Glu Cys
 55 65 70 75 80

Asp Cys Pro Pro Asn Phe Pro Thr Ala Met Tyr Cys Asp Asn Arg Asn
 60 85 90 95

Leu Lys Tyr Leu Pro Phe Val Pro Ser Arg Met Lys Tyr Val Tyr Phe
 65 100 105 110

Gln Asn Asn Gln Ile Thr Ser Ile Gln Glu Gly Val Phe Asp Asn Ala
 70 115 120 125

Thr Gly Leu Leu Trp Ile Ala Leu His Gly Asn Gln Ile Thr Ser Asp
 130 135 140

5 Lys Val Gly Arg Lys Val Phe Ser Lys Leu Arg His Leu Glu Arg Leu
 145 150 155 160

10 Tyr Leu Asp His Asn Asn Leu Thr Arg Met Pro Gly Pro Leu Pro Arg
 165 170 175

15 Ser Leu Arg Glu Leu His Leu Asp His Asn Gln Ile Ser Arg Val Pro
 180 185 190

20 Asn Asn Ala Leu Glu Gly Leu Glu Asn Leu Thr Ala Leu Tyr Leu Gln
 195 200 205

25 His Asn Glu Ile Gln Glu Val Gly Ser Ser Met Arg Gly Leu Arg Ser
 210 215 220

30 Leu Tyr Leu Leu Asp Leu Ser Tyr Asn His Leu Arg Lys Val Pro Asp
 225 230 235 240

35 Gly Leu Pro Ser Ala Leu Glu Gln Leu Tyr Met Glu His Asn Asn Val
 245 250 255

40 Tyr Thr Val Pro Asp Ser Tyr Phe Arg Gly Ala Pro Lys Leu Leu Tyr
 260 265 270

45 Val Arg Leu Ser His Asn Ser Leu Thr Asn Asn Gly Leu Ala Ser Asn
 275 280 285

50 Thr Phe Asn Ser Ser Leu Leu Glu Leu Asp Leu Ser Tyr Asn Gln
 290 295 300

55 Leu Gln Lys Ile Pro Pro Val Asn Thr Asn Leu Glu Asn Leu Tyr Leu
 305 310 315 320

60 Gln Gly Asn Arg Ile Asn Glu Phe Ser Ile Ser Ser Phe Cys Thr Val
 325 330 335

65 Val Asp Val Val Asn Phe Ser Gln Leu Gln Val Val Arg Leu Asp Gly
 340 345 350

70 Asn Glu Met Lys Arg Ser Ala Met Pro Ala Glu Ala Pro Leu Cys Leu
 355 360 365

Arg Leu Ala Ser Leu Ile Glu Ile
 370 375

65 <210> 131
 <211> 897
 <212> PRT
 <213> Homo sapiens

<300>
<308> Swissprot/P32927
<309> 1993-10-01
<313> (1)..(897)

5

<400> 131

Met Val Leu Ala Gln Gly Leu Leu Ser Met Ala Leu Leu Ala Leu Cys
1 5 10 15

Trp Glu Arg Ser Leu Ala Gly Ala Glu Glu Thr Ile Pro Leu Gln Thr
20 25 30

Leu Arg Cys Tyr Asn Asp Tyr Thr Ser His Ile Thr Cys Arg Trp Ala
35 40 45

Asp Thr Gln Asp Ala Gln Arg Leu Val Asn Val Thr Leu Ile Arg Arg
50 55 60

Val Asn Glu Asp Leu Leu Glu Pro Val Ser Cys Asp Leu Ser Asp Asp
65 70 75 80

Met Pro Trp Ser Ala Cys Pro His Pro Arg Cys Val Pro Arg Arg Cys
85 90 95

Val Ile Pro Cys Gln Ser Phe Val Val Thr Asp Val Asp Tyr Phe Ser
100 105 110

Phe Gln Pro Asp Arg Pro Leu Gly Thr Arg Leu Thr Val Thr Leu Thr
115 120 125

Gln His Val Gln Pro Pro Glu Pro Arg Asp Leu Gln Ile Ser Thr Asp
130 135 140

Gln Asp His Phe Leu Leu Thr Trp Ser Val Ala Leu Gly Ser Pro Gln
145 150 155 160

Ser His Trp Leu Ser Pro Gly Asp Leu Glu Phe Glu Val Val Tyr Lys
165 170 175

Arg Leu Gln Asp Ser Trp Glu Asp Ala Ala Ile Leu Leu Ser Asn Thr
180 185 190

Ser Gln Ala Thr Leu Gly Pro Glu His Leu Met Pro Ser Ser Thr Tyr
195 200 205

Val Ala Arg Val Arg Thr Arg Leu Ala Pro Gly Ser Arg Leu Ser Gly
210 215 220

Arg Pro Ser Lys Trp Ser Pro Glu Val Cys Trp Asp Ser Gln Pro Gly
225 230 235 240

Asp Glu Ala Gln Pro Gln Asn Leu Glu Cys Phe Phe Asp Gly Ala Ala
245 250 255

70

val Leu Ser Cys Ser Trp Glu Val Arg Lys Glu Val Ala Ser Ser Val
260 265 270

5 Ser Phe Gly Leu Phe Tyr Lys Pro Ser Pro Asp Ala Gly Glu Glu Glu
275 280 285

10 Cys Ser Pro Val Leu Arg Glu Gly Leu Gly Ser Leu His Thr Arg His
290 295 300

15 His Cys Gln Ile Pro Val Pro Asp Pro Ala Thr His Gly Gln Tyr Ile
305 310 315 320

20 val Ser Val Gln Pro Arg Arg Ala Glu Lys His Ile Lys Ser Ser Val
325 330 335

Asn Ile Gln Met Ala Pro Pro Ser Leu Asn Val Thr Lys Asp Gly Asp
340 345 350

25 Ser Tyr Ser Leu Arg Trp Glu Thr Met Lys Met Arg Tyr Glu His Ile
355 360 365

30 Asp His Thr Phe Glu Ile Gln Tyr Arg Lys Asp Thr Ala Thr Trp Lys
370 375 380

35 Asp Ser Lys Thr Glu Thr Leu Gln Asn Ala His Ser Met Ala Leu Pro
385 390 395 400

40 Ala Leu Glu Pro Ser Thr Arg Tyr Trp Ala Arg Val Arg Val Arg Thr
405 410 415

45 Ser Arg Thr Gly Tyr Asn Gly Ile Trp Ser Glu Trp Ser Glu Ala Arg
420 425 430

50 Ser Trp Asp Thr Glu Ser Val Leu Pro Met Trp Val Leu Ala Leu Ile
435 440 445

55 val Ile Phe Leu Thr Ile Ala Val Leu Leu Ala Leu Arg Phe Cys Gly
450 455 460

Ile Tyr Gly Tyr Arg Leu Arg Arg Lys Trp Glu Glu Lys Ile Pro Asn
465 470 475 480

60 Pro Ser Lys Ser His Leu Phe Gln Asn Gly Ser Ala Glu Leu Trp Pro
485 490 495

Pro Gly Ser Met Ser Ala Phe Thr Ser Gly Ser Pro Pro His Gln Gly
500 505 510

65 Pro Trp Gly Ser Arg Phe Pro Glu Leu Glu Gly Val Phe Pro Val Gly
515 520 525

70 Phe Gly Asp Ser Glu Val Ser Pro Leu Thr Ile Glu Asp Pro Lys His

	530	535	540
5	Val Cys Asp Pro Pro Ser Gly Pro Asp Thr Thr Pro Ala Ala Ser Asp		
	545 550		555 560
10	Leu Pro Thr Glu Gln Pro Pro Ser Pro Gln Pro Gly Pro Pro Ala Ala		
	565 570		575 580
15	Ser His Thr Pro Glu Lys Gln Ala Ser Ser Phe Asp Phe Asn Gly Pro		
	580 585		590 595
20	Tyr Leu Gly Pro Pro His Ser Arg Ser Leu Pro Asp Ile Leu Gly Gln		
	595 600		605 610
25	Pro Glu Pro Pro Gln Glu Gly Gly Ser Gln Lys Ser Pro Pro Pro Gly		
	610 615		620 625
30	Ser Leu Glu Tyr Leu Cys Leu Pro Ala Gly Gly Gln Val Gln Leu Val		
	625 630		635 640
35	Pro Leu Ala Gln Ala Met Gly Pro Gly Gln Ala Val Glu Val Glu Arg		
	645 650		655 660
40	Arg Pro Ser Gln Gly Ala Ala Gly Ser Pro Ser Leu Glu Ser Gly Gly		
	660 665		670 675
45	Gly Pro Ala Pro Pro Ala Leu Gly Pro Arg Val Gly Gly Gln Asp Gln		
	675 680		685 690
50	Lys Asp Ser Pro Val Ala Ile Pro Met Ser Ser Gly Asp Thr Glu Asp		
	690 695		700 705
55	Pro Gly Val Ala Ser Gly Tyr Val Ser Ser Ala Asp Leu Val Phe Thr		
	710 715		720 725
60	Pro Asn Ser Gly Ala Ser Ser Val Ser Leu Val Pro Ser Leu Gly Leu		
	730 735		740 745
65	Pro Ser Asp Gln Thr Pro Ser Leu Cys Pro Gly Leu Ala Ser Gly Pro		
	750 755		760 765
70	Pro Gly Ala Pro Gly Pro Val Lys Ser Gly Phe Glu Gly Tyr Val Glu		
	770 775		780 785
	Leu Pro Pro Ile Glu Gly Arg Ser Pro Arg Ser Pro Arg Asn Asn Pro		
	790 795		800 805
	Ala Asp Val Ser Pro Thr Ser Pro Gln Pro Glu Gly Leu Leu Val Leu		
	810 815		

Gln Gln Val Gly Asp Tyr Cys Phe Leu Pro Gly Leu Gly Pro Gly Pro
 820 825 830

5 Leu Ser Leu Arg Ser Lys Pro Ser Ser Pro Gly Pro Gly Pro Glu Ile
 835 840 845

10 Lys Asn Leu Asp Gln Ala Phe Gln Val Lys Lys Pro Pro Gly Gln Ala
 850 855 860

15 Val Pro Gln Val Pro Val Ile Gln Leu Phe Lys Ala Leu Lys Gln Gln
 865 870 875 880

20 Asp Tyr Leu Ser Leu Pro Pro Trp Glu Val Asn Lys Pro Gly Glu Val
 885 890 895

Cys

25 <210> 132
 <211> 261
 <212> PRT
 30 <213> Homo sapiens

35 <300>
 <308> Swissprot/Q60493
 <309> 1996-11-01
 <313> (1)..(261)

40 <400> 132

Met Ile Tyr Lys Cys Pro Met Cys Arg Glu Phe Phe Ser Glu Arg Ala
 1 5 10 15

45 Asp Leu Phe Met His Gln Lys Val His Thr Ala Glu Lys Pro His Lys
 20 25 30

50 Cys Asp Lys Cys Asp Lys Gly Phe Phe His Ile Ser Glu Leu His Ile
 35 40 45

55 His Trp Arg Asp His Thr Gly Glu Lys Val Tyr Lys Cys Asp Asp Cys
 50 55 60

60 Gly Lys Asp Phe Ser Thr Thr Lys Leu Asn Arg His Lys Lys Ile
 65 70 75 80

65 His Thr Val Glu Lys Pro Tyr Lys Cys Tyr Glu Cys Gly Lys Ala Phe
 85 90 95

70 Asn Trp Ser Pro His Leu Gln Ile His Met Arg Val His Thr Gly Glu
 100 105 110

75 Lys Pro Tyr Val Cys Ser Glu Cys Gly Arg Gly Phe Ser Asn Ser Ser
 115 120 125

80 Asn Leu Cys Met His Gln Arg Val His Thr Gly Glu Lys Pro Phe Lys

	130	135	140
5	Cys Glu Glu Cys Gly Lys Ala Phe Arg His Thr Ser Ser Leu Cys Met 145 150 155 160		
10	His Gln Arg Val His Thr Gly Glu Lys Pro Tyr Lys Cys Tyr Glu Cys 165 170 175		
15	Gly Lys Ala Phe Ser Gln Ser Ser Ser Leu Cys Ile His Gln Arg Val 180 185 190		
20	His Thr Gly Glu Lys Pro Tyr Arg Cys Cys Gly Cys Gly Lys Ala Phe 195 200 205		
25	Ser Gln Ser Ser Ser Leu Cys Ile His Gln Arg Val His Thr Gly Glu 210 215 220	Lys Pro Phe Lys Cys Asp Glu Cys Gly Lys Ala Phe Ser Gln Ser Thr 225 230 235 240	
30	Ser Leu Cys Ile His Gln Arg Val His Thr Lys Glu Arg Asn His Leu 245 250 255		
	Lys Ile Ser Val Ile 260		
35	<210> 133 <211> 296 <212> PRT <213> Homo sapiens		
40	<300> <308> Swissprot/P04233 <309> 1987-03-20 <313> (1)..(296)		
45	<400> 133		
50	Met His Arg Arg Arg Ser Arg Ser Cys Arg Glu Asp Gln Lys Pro Val 1 5 10 15	Met Asp Asp Gln Arg Asp Leu Ile Ser Asn Asn Glu Gln Leu Pro Met 20 25 30	
55	Leu Gly Arg Arg Pro Gly Ala Pro Glu Ser Lys Cys Ser Arg Gly Ala 35 40 45		
60	Leu Tyr Thr Gly Phe Ser Ile Leu Val Thr Leu Leu Leu Ala Gly Gln 50 55 60		
65	Ala Thr Thr Ala Tyr Phe Leu Tyr Gln Gln Gln Gly Arg Leu Asp Lys 65 70 75 80		
70	Leu Thr Val Thr Ser Gln Asn Leu Gln Leu Glu Asn Leu Arg Met Lys 85 90 95		

Leu Pro Lys Pro Pro Lys Pro Val Ser Lys Met Arg Met Ala Thr Pro
 100 105 110

5 Leu Leu Met Gln Ala Leu Pro Met Gly Ala Leu Pro Gln Gly Pro Met
 115 120 125

10 Gln Asn Ala Thr Lys Tyr Gly Asn Met Thr Glu Asp His Val Met His
 130 135 140

15 Leu Leu Gln Asn Ala Asp Pro Leu Lys Val Tyr Pro Pro Leu Lys Gly
 145 150 155 160

20 Ser Phe Pro Glu Asn Leu Arg His Leu Lys Asn Thr Met Glu Thr Ile
 165 170 175

25 Asp Trp Lys Val Phe Glu Ser Trp Met His His Trp Leu Leu Phe Glu
 180 185 190

30 Met Ser Arg His Ser Leu Glu Gln Lys Pro Thr Asp Ala Pro Pro Lys
 195 200 205

35 Val Leu Thr Lys Cys Gln Glu Glu Val Ser His Ile Pro Ala Val His
 210 215 220

40 Pro Gly Ser Phe Arg Pro Lys Cys Asp Glu Asn Gly Asn Tyr Leu Pro
 225 230 235 240

Leu Gln Cys Tyr Gly Ser Ile Gly Tyr Cys Trp Cys Val Phe Pro Asn
 245 250 255

45 Gly Thr Glu Val Pro Asn Thr Arg Ser Arg Gly His His Asn Cys Ser
 260 265 270

50 Glu Ser Leu Glu Leu Glu Asp Pro Ser Ser Gly Leu Gly Val Thr Lys
 275 280 285

55 Gln Asp Leu Gly Pro Val Pro Met
 290 295

<210> 134
 <211> 163
 <212> PRT
 <213> Homo sapiens

60 <300>
 <308> Swissprot/Q99969
 <309> 2000-05-30
 <313> (1)..(163)

65 <400> 134
 Met Arg Arg Leu Leu Ile Pro Leu Ala Leu Trp Leu Gly Ala Val Gly
 1 5 10 15

70 Val Gly Val Ala Glu Leu Thr Glu Ala Gln Arg Arg Gly Leu Gln Val

83

20

25

30

Ala Leu Glu Glu Phe His His Pro Pro Val Gln Trp Ala Phe Gln
 5 35 40 45

Glu Thr Ser Val Glu Ser Ala Val Asp Thr Pro Phe Pro Ala Gly Ile
 10 50 55 60

Phe Val Arg Leu Glu Phe Lys Leu Gln Gln Thr Ser Cys Arg Lys Arg
 15 65 70 75 80

Asp Trp Lys Lys Pro Glu Cys Lys Val Arg Pro Asn Gly Arg Lys Arg
 20 85 90 95

Lys Cys Leu Ala Cys Ile Lys Leu Gly Ser Glu Asp Lys Val Leu Gly
 25 100 105 110

Arg Leu Val His Cys Pro Ile Glu Thr Gln Val Leu Arg Glu Ala Glu
 30 115 120 125

Glu His Gln Glu Thr Gln Cys Leu Arg Val Gln Arg Ala Gly Glu Asp
 35 130 135 140

Pro His Ser Phe Tyr Phe Pro Gly Gln Phe Ala Phe Ser Lys Ala Leu
 40 145 150 155 160

Pro Arg Ser

45 <210> 135
 <211> 2386
 <212> PRT
 <213> Homo sapiens

50 <300>
 <308> Swissprot/P02751
 <309> 1986-07-21
 <313> (1)..(2386)

55 <400> 135

Met Leu Arg Gly Pro Gly Pro Gly Leu Leu Leu Leu Ala Val Gln Cys
 1 5 10 15

60 Leu Gly Thr Ala Val Pro Ser Thr Gly Ala Ser Lys Ser Lys Arg Gln
 20 25 30

Ala Gln Gln Met Val Gln Pro Gln Ser Pro Val Ala Val Ser Gln Ser
 65 35 40 45

Lys Pro Gly Cys Tyr Asp Asn Gly Lys His Tyr Gln Ile Asn Gln Gln
 70 50 55 60 65

Trp Glu Arg Thr Tyr Leu Gly Asn Ala Leu Val Cys Thr Cys Tyr Gly
 75 80

Gly Ser Arg Gly Phe Asn Cys Glu Ser Lys Pro Glu Ala Glu Glu Thr
 85 80 90 95

5 Cys Phe Asp Lys Tyr Thr Gly Asn Thr Tyr Arg Val Gly Asp Thr Tyr
 100 105 110

10 Glu Arg Pro Lys Asp Ser Met Ile Trp Asp Cys Thr Cys Ile Gly Ala
 115 120 125

15 Gly Arg Gly Arg Ile Ser Cys Thr Ile Ala Asn Arg Cys His Glu Gly
 130 135 140

20 Gly Gln Ser Tyr Lys Ile Gly Asp Thr Trp Arg Arg Pro His Glu Thr
 145 150 155 160

25 Gly Gly Tyr Met Leu Glu Cys Val Cys Leu Gly Asn Gly Lys Gly Glu
 165 170 175

30 Trp Thr Cys Lys Pro Ile Ala Glu Lys Cys Phe Asp His Ala Ala Gly
 180 185 190

35 Thr Ser Tyr Val Val Gly Glu Thr Trp Glu Lys Pro Tyr Gln Gly Trp
 195 200 205

40 Met Met Val Asp Cys Thr Cys Leu Gly Glu Gly Ser Gly Arg Ile Thr
 210 215 220

45 Cys Thr Ser Arg Asn Arg Cys Asn Asp Gln Asp Thr Arg Thr Ser Tyr
 225 230 235 240

50 Arg Ile Gly Asp Thr Trp Ser Lys Lys Asp Asn Arg Gly Asn Leu Leu
 245 250 255

55 Gln Cys Ile Cys Thr Gly Asn Gly Arg Gly Glu Trp Lys Cys Glu Arg
 260 265 270

60 His Thr Ser Val Gln Thr Thr Ser Ser Gly Ser Gly Pro Phe Thr Asp
 275 280 285

65 Val Arg Ala Ala Val Tyr Gln Pro Gln Pro His Pro Gln Pro Pro Pro
 290 295 300

70 Tyr Gly His Cys Val Thr Asp Ser Gly Val Val Tyr Ser Val Gly Met
 305 310 315 320

75 Gln Trp Leu Lys Thr Gln Gly Asn Lys Gln Met Leu Cys Thr Cys Leu
 325 330 335

80 Gly Asn Gly Val Ser Cys Gln Glu Thr Ala Val Thr Gln Thr Tyr Gly
 340 345 350

85 Gly Asn Ser Asn Gly Glu Pro Cys Val Leu Pro Phe Thr Tyr Asn Gly

85

355

360

365

5 Arg Thr Phe Tyr Ser Cys Thr Thr Glu Gly Arg Gln Asp Gly His Leu
370 375 380

10 Trp Cys Ser Thr Thr Ser Asn Tyr Glu Gln Asp Gln Lys Tyr Ser Phe
385 390 395 400

15 Cys Thr Asp His Thr Val Leu Val Gln Thr Gln Gly Gly Asn Ser Asn
405 410 415

20 Gly Ala Leu Cys His Phe Pro Phe Leu Tyr Asn Asn His Asn Tyr Thr
420 425 430

25 Asp Cys Thr Ser Glu Gly Arg Arg Asp Asn Met Lys Trp Cys Gly Thr
435 440 445

30 Thr Gln Asn Tyr Asp Ala Asp Gln Lys Phe Gly Phe Cys Pro Met Ala
450 455 460

35 Ala His Glu Glu Ile Cys Thr Thr Asn Glu Gly Val Met Tyr Arg Ile
465 470 475 480

40 Gly Asp Gln Trp Asp Lys Gln His Asp Met Gly His Met Met Arg Cys
485 490 495

45 Thr Cys Val Gly Asn Gly Arg Gly Glu Trp Thr Cys Ile Ala Tyr Ser
500 505 510

50 Gln Leu Arg Asp Gln Cys Ile Val Asp Asp Ile Thr Tyr Asn Val Asn
515 520 525

55 Asp Thr Phe His Lys Arg His Glu Glu Gly His Met Leu Asn Cys Thr
530 535 540

60 Cys Phe Gly Gln Gly Arg Gly Arg Trp Lys Cys Asp Pro Val Asp Gln
545 550 555 560

65 Cys Gln Asp Ser Glu Thr Gly Thr Phe Tyr Gln Ile Gly Asp Ser Trp
565 570 575

70 Glu Lys Tyr Val His Gly Val Arg Tyr Gln Cys Tyr Cys Tyr Gly Arg
580 585 590

75 Gly Ile Gly Glu Trp His Cys Gln Pro Leu Gln Thr Tyr Pro Ser Ser
595 600 605

80 Ser Gly Pro Val Glu Val Phe Ile Thr Glu Thr Pro Ser Gln Pro Asn
610 615 620

85 Ser His Pro Ile Gln Trp Asn Ala Pro Gln Pro Ser His Ile Ser Lys
625 630 635 640

Tyr Ile Leu Arg Trp Arg Pro Lys Asn Ser Val Gly Arg Trp Lys Glu
 645 650 655

5 Ala Thr Ile Pro Gly His Leu Asn Ser Tyr Thr Ile Lys Gly Leu Lys
 660 665 670

10 Pro Gly Val Val Tyr Glu Gly Gln Leu Ile Ser Ile Gln Gln Tyr Gly
 675 680 685

15 His Gln Glu Val Thr Arg Phe Asp Phe Thr Thr Ser Thr Ser Thr
 690 695 700

20 Pro Val Thr Ser Asn Thr Val Thr Gly Glu Thr Thr Pro Phe Ser Pro
 705 710 715 720

25 Leu Val Ala Thr Ser Glu Ser Val Thr Glu Ile Thr Ala Ser Ser Phe
 725 730 735

30 Val Val Ser Trp Val Ser Ala Ser Asp Thr Val Ser Gly Phe Arg Val
 740 745 750

35 Glu Tyr Glu Leu Ser Glu Glu Gly Asp Glu Pro Gln Tyr Leu Asp Leu
 755 760 765

40 Pro Ser Thr Ala Thr Ser Val Asn Ile Pro Asp Leu Leu Pro Gly Arg
 770 775 780

45 Lys Tyr Ile Val Asn Val Tyr Gln Ile Ser Glu Asp Gly Glu Gln Ser
 785 790 795 800

50 Leu Ile Leu Ser Thr Ser Gln Thr Thr Ala Pro Asp Ala Pro Pro Asp
 805 810 815

55 Pro Thr Val Asp Gln Val Asp Asp Thr Ser Ile Val Val Arg Trp Ser
 820 825 830

60 Arg Pro Gln Ala Pro Ile Thr Gly Tyr Arg Ile Val Tyr Ser Pro Ser
 835 840 845

65 Val Glu Gly Ser Ser Thr Glu Leu Asn Leu Pro Glu Thr Ala Asn Ser
 850 855 860

70 Val Thr Leu Ser Asp Leu Gln Pro Gly Val Gln Tyr Asn Ile Thr Ile
 865 870 875 880

75 Tyr Ala Val Glu Glu Asn Gln Glu Ser Thr Pro Val Val Ile Gln Gln
 885 890 895

80 Glu Thr Thr Gly Thr Pro Arg Ser Asp Thr Val Pro Ser Pro Arg Asp
 900 905 910

85 Leu Gln Phe Val Glu Val Thr Asp Val Lys Val Thr Ile Met Trp Thr

	915	920	925
5	Pro Pro Glu Ser Ala Val Thr Gly Tyr Arg Val Asp Val Ile Pro Val 930 935 940 940		
10	Asn Leu Pro Gly Glu His Gly Gln Arg Leu Pro Ile Ser Arg Asn Thr 945 950 955 955 960		
15	Phe Ala Glu Val Thr Gly Leu Ser Pro Gly Val Thr Tyr Tyr Phe Lys 965 970 975		
20	Val Phe Ala Val Ser His Gly Arg Glu Ser Lys Pro Leu Thr Ala Gln 980 985 990		
25	Gln Thr Thr Lys Leu Asp Ala Pro Thr Asn Leu Gln Phe Val Asn Glu 995 1000 1005		
30	Thr Asp Ser Thr Val Leu Val Arg Trp Thr Pro Pro Arg Ala Gln 1010 1015 1020		
35	Ile Thr Gly Tyr Arg Leu Thr Val Gly Leu Thr Arg Arg Gly Gln 1025 1030 1035		
40	Pro Arg Gln Tyr Asn Val Gly Pro Ser Val Ser Lys Tyr Pro Leu 1040 1045 1050		
45	Arg Asn Leu Gln Pro Ala Ser Glu Tyr Thr Val Ser Leu Val Ala 1055 1060 1065		
50	Ile Lys Gly Asn Gln Glu Ser Pro Lys Ala Thr Gly Val Phe Thr 1070 1075 1080		
55	Thr Leu Gln Pro Gly Ser Ser Ile Pro Pro Tyr Asn Thr Glu Val 1085 1090 1095		
60	Thr Glu Thr Thr Ile Val Ile Thr Trp Thr Pro Ala Pro Arg Ile 1100 1105 1110		
65	Gly Phe Lys Leu Gly Val Arg Pro Ser Gln Gly Gly Glu Ala Pro 1115 1120 1125		
70	Arg Glu Val Thr Ser Asp Ser Gly Ser Ile Val Val Ser Gly Leu 1130 1135 1140		
	Thr Pro Gly Val Glu Tyr Val Tyr Thr Ile Gln Val Leu Arg Asp 1145 1150 1155		
	Gly Gln Glu Arg Asp Ala Pro Ile Val Asn Lys Val Val Thr Pro 1160 1165 1170		
	Leu Ser Pro Pro Thr Asn Leu His Leu Glu Ala Asn Pro Asp Thr 1175 1180 1185		

Gly Val Leu Thr Val Ser Trp Glu Arg Ser Thr Thr Pro Asp Ile
 1190 1195 1200
 5 Thr Gly Tyr Arg Ile Thr Thr Pro Thr Asn Gly Gln Gln Gly
 1205 1210 1215
 10 Asn Ser Leu Glu Glu Val Val His Ala Asp Gln Ser Ser Cys Thr
 1220 1225 1230
 15 Phe Asp Asn Leu Ser Pro Gly Leu Glu Tyr Asn Val Ser Val Tyr
 1235 1240 1245
 20 Thr Val Lys Asp Asp Lys Glu Ser Val Pro Ile Ser Asp Thr Ile
 1250 1255 1260
 25 Ile Pro Ala Val Pro Pro Thr Asp Leu Arg Phe Thr Asn Ile
 1265 1270 1275
 30 Gly Pro Asp Thr Met Arg Val Thr Trp Ala Pro Pro Ser Ile
 1280 1285 1290
 35 Asp Leu Thr Asn Phe Leu Val Arg Tyr Ser Pro Val Lys Asn Glu
 1295 1300 1305
 40 Glu Asp Val Ala Glu Leu Ser Ile Ser Pro Ser Asp Asn Ala Val
 1310 1315 1320
 45 Val Leu Thr Asn Leu Leu Pro Gly Thr Glu Tyr Val Val Ser Val
 1325 1330 1335
 50 Ser Ser Val Tyr Glu Gln His Glu Ser Thr Pro Leu Arg Gly Arg
 1340 1345 1350
 55 Gln Lys Thr Gly Leu Asp Ser Pro Thr Gly Ile Asp Phe Ser Asp
 1355 1360 1365
 60 Ile Thr Ala Asn Ser Phe Thr Val His Trp Ile Ala Pro Arg Ala
 1370 1375 1380
 65 Thr Ile Thr Gly Tyr Arg Ile Arg His His Pro Glu His Phe Ser
 1385 1390 1395
 70 Gly Arg Pro Arg Glu Asp Arg Val Pro His Ser Arg Asn Ser Ile
 1400 1405 1410
 75 Thr Leu Thr Asn Leu Thr Pro Gly Thr Glu Tyr Val Val Ser Ile
 1415 1420 1425
 80 Val Ala Leu Asn Gly Arg Glu Glu Ser Pro Leu Leu Ile Gly Gln
 1430 1435 1440
 85 Gln Ser Thr Val Ser Asp Val Pro Arg Asp Leu Glu Val Val Ala

	1445	1450	1455	
5	Ala Thr Pro Thr Ser Leu Leu Ile Ser Trp Asp Ala	1460 1465	1470	Pro Ala Val
10	Thr Val Arg Tyr Tyr Arg Ile Thr Tyr Gly Glu Thr	1475 1480	1485	Gly Gly Asn
	Ser Pro Val Gln Glu Phe Thr Val Pro Gly Ser Lys	1490 1495	1500	Ser Thr Ala
15	Thr Ile Ser Gly Leu Lys Pro Gly Val Asp Tyr Thr	1505 1510	1515	Ile Thr Val
20	Tyr Ala Val Thr Gly Arg Gly Asp Ser Pro Ala Ser	1520 1525	1530	Ser Lys Pro
25	Ile Ser Ile Asn Tyr Arg Thr Glu Ile Asp Lys Pro	1535 1540	1545	Ser Gln Met
30	Gln Val Thr Asp Val Gln Asp Asn Ser Ile Ser Val	1550 1555	1560	Lys Trp Leu
	Pro Ser Ser Ser Pro Val Thr Gly Tyr Arg Val Thr	1565 1570	1575	Thr Thr Pro
35	Lys Asn Gly Pro Gly Pro Thr Lys Thr Lys Thr Ala	1580 1585	1590	Gly Pro Asp
40	Gln Thr Glu Met Thr Ile Glu Gly Leu Gln Pro Thr	1595 1600	1605	Val Glu Tyr
45	Val Val Ser Val Tyr Ala Gln Asn Pro Ser Gly Glu	1610 1615	1620	Ser Gln Pro
50	Leu Val Gln Thr Ala Val Thr Asn Ile Asp Arg Pro	1625 1630	1635	Lys Gly Leu
	Ala Phe Thr Asp Val Asp Val Asp Ser Ile Lys Ile	1640 1645	1650	Ala Trp Glu
55	Ser Pro Gln Gly Gln Val Ser Arg Tyr Arg Val Thr	1655 1660	1665	Tyr Ser Ser
60	Pro Glu Asp Gly Ile His Glu Leu Phe Pro Ala Pro	1670 1675	1680	Asp Gly Glu
65	Glu Asp Thr Ala Glu Leu Gln Gly Leu Arg Pro Gly	1685 1690	1695	Ser Glu Tyr
70	Thr Val Ser Val Val Ala Leu His Asp Asp Met Glu	1700 1705	1710	Ser Gln Pro

Leu Ile Gly Thr Gln Ser Thr Ala Ile Pro Ala Pro Thr Asp Leu
 1715 1720 1725

5 Lys Phe Thr Gln Val Thr Pro Thr Ser Leu Ser Ala Gln Trp Thr
 1730 1735 1740

10 Pro Pro Asn Val Gln Leu Thr Gly Tyr Arg Val Arg Val Thr Pro
 1745 1750 1755

15 Lys Glu Lys Thr Gly Pro Met Lys Glu Ile Asn Leu Ala Pro Asp
 1760 1765 1770

20 Ser Ser Val Val Val Ser Gly Leu Met Val Ala Thr Lys Tyr
 1775 1780 1785

25 Glu Val Ser Val Tyr Ala Leu Lys Asp Thr Leu Thr Ser Arg Pro
 1790 1795 1800

30 Ala Gln Gly Val Val Thr Thr Leu Glu Asn Val Ser Pro Pro Arg
 1805 1810 1815

35 Arg Ala Arg Val Thr Asp Ala Thr Glu Thr Thr Ile Thr Ile Ser
 1820 1825 1830

40 Trp Arg Thr Lys Thr Glu Thr Ile Thr Gly Phe Gln Val Asp Ala
 1835 1840 1845

45 Val Pro Ala Asn Gly Gln Thr Pro Ile Gln Arg Thr Ile Lys Pro
 1850 1855 1860

50 Asp Val Arg Ser Tyr Thr Ile Thr Gly Leu Gln Pro Gly Thr Asp
 1865 1870 1875

55 Tyr Lys Ile Tyr Leu Tyr Thr Leu Asn Asp Asn Ala Arg Ser Ser
 1880 1885 1890

60 Pro Val Val Ile Asp Ala Ser Thr Ala Ile Asp Ala Pro Ser Asn
 1895 1900 1905

65 Leu Arg Phe Leu Ala Thr Thr Pro Asn Ser Leu Leu Val Ser Trp
 1910 1915 1920

70 Gln Pro Pro Arg Ala Arg Ile Thr Gly Tyr Ile Ile Lys Tyr Glu
 1925 1930 1935

75 Lys Pro Gly Ser Pro Pro Arg Glu Val Val Pro Arg Pro Arg Pro
 1940 1945 1950

80 Gly Val Thr Glu Ala Thr Ile Thr Gly Leu Glu Pro Gly Thr Glu
 1955 1960 1965

85 Tyr Thr Ile Tyr Val Ile Ala Leu Lys Asn Asn Gln Lys Ser Glu

	1970	1975	1980
5	Pro Leu Ile Gly Arg Lys Lys	Thr Asp Glu Leu Pro Gln Leu Val	
	1985 1990	1995	
10	Thr Leu Pro His Pro Asn Leu His Gly Pro Glu Ile Leu Asp Val		
	2000 2005	2010	
15	Pro Ser Thr Val Gln Lys Thr Pro Phe Val Thr His Pro Gly Tyr		
	2015 2020	2025	
20	Asp Thr Gly Asn Gly Ile Gln Leu Pro Gly Thr Ser Gly Gln Gln		
	2030 2035	2040	
25	Pro Ser Val Gly Gln Gln Met Ile Phe Glu Glu His Gly Phe Arg		
	2045 2050	2055	
30	Arg Thr Thr Pro Pro Thr Thr Ala Thr Pro Ile Arg His Arg Pro		
	2060 2065	2070	
35	Arg Pro Tyr Pro Pro Asn Val Gly Glu Glu Ile Gln Ile Gly His		
	2075 2080	2085	
40	Ile Pro Arg Glu Asp Val Asp Tyr His Leu Tyr Pro His Gly Pro		
	2090 2095	2100	
45	Gly Leu Asn Pro Asn Ala Ser Thr Gly Gln Glu Ala Leu Ser Gln		
	2105 2110	2115	
50	Thr Thr Ile Ser Trp Ala Pro Phe Gln Asp Thr Ser Glu Tyr Ile		
	2120 2125	2130	
55	Ile Ser Cys His Pro Val Gly Thr Asp Glu Glu Pro Leu Gln Phe		
	2135 2140	2145	
60	Arg Val Pro Gly Thr Ser Thr Ser Ala Thr Leu Thr Gly Leu Thr		
	2150 2155	2160	
65	Arg Gly Ala Thr Tyr Asn Ile Ile Val Glu Ala Leu Lys Asp Gln		
	2165 2170	2175	
70	Gln Arg His Lys Val Arg Glu Glu Val Val Thr Val Gly Asn Ser		
	2180 2185	2190	
	Val Asn Glu Gly Leu Asn Gln Pro Thr Asp Asp Ser Cys Phe Asp		
	2195 2200	2205	
	Pro Tyr Thr Val Ser His Tyr Ala Val Gly Asp Glu Trp Glu Arg		
	2210 2215	2220	
	Met Ser Glu Ser Gly Phe Lys Leu Leu Cys Gln Cys Leu Gly Phe		
	2225 2230	2235	

Gly Ser Gly His Phe Arg Cys Asp Ser Ser Arg Trp Cys His Asp
 2240 2245 2250 2255 2260 2265 2270 2275 2280 2285 2290 2295 2300 2305 2310 2315 2320 2325 2330 2335 2340 2345 2350 2355 2360 2365 2370 2375 2380 2385
 5 Asn Gly Val Asn Tyr Lys Ile Gly Glu Lys Trp Asp Arg Gln Gly
 10 Glu Asn Gly Gln Met Met Ser Cys Thr Cys Leu Gly Asn Gly Lys
 15 Gly Glu Phe Lys Cys Asp Pro His Glu Ala Thr Cys Tyr Asp Asp
 20 Gly Lys Thr Tyr His Val Gly Glu Gln Trp Gln Lys Glu Tyr Leu
 25 Gly Ala Ile Cys Ser Cys Thr Cys Phe Gly Gly Gln Arg Gly Trp
 30 Arg Cys Asp Asn Cys Arg Arg Pro Gly Gly Glu Pro Ser Pro Glu
 35 Gly Thr Thr Gly Gln Ser Tyr Asn Gln Tyr Ser Gln Arg Tyr His
 40 Gln Arg Thr Asn Thr Asn Val Asn Cys Pro Ile Glu Cys Phe Met
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 45 <213> Homo sapiens
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 <309> 1988-08-01
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 <400> 136
 55 Met Trp Gln Leu Trp Ala Ser Leu Cys Cys Leu Leu Val Leu Ala Asn
 1 5 10 15
 60 Ala Arg Ser Arg Pro Ser Phe His Pro Val Ser Asp Glu Leu Val Asn
 65 Tyr Val Asn Lys Arg Asn Thr Thr Trp Gln Ala Gly His Asn Phe Tyr
 70 Asn Val Asp Met Ser Tyr Leu Lys Arg Leu Cys Gly Thr Phe Leu Gly
 70 Gly Pro Lys Pro Pro Gln Arg Val Met Phe Thr Glu Asp Leu Lys Leu

65 70 75 80

5 Pro Ala Ser Phe Asp Ala Arg Glu Gln Trp Pro Gln Cys Pro Thr Ile
85 90 95

10 Lys Glu Ile Arg Asp Gln Gly Ser Cys Gly Ser Cys Trp Ala Phe Gly
100 105 110

15 Ala Val Glu Ala Ile Ser Asp Arg Ile Cys Ile His Thr Asn Ala His
115 120 125

20 Val Ser Val Glu Val Ser Ala Glu Asp Leu Leu Thr Cys Cys Gly Ser
130 135 140

25 Met Cys Gly Asp Gly Cys Asn Gly Gly Tyr Pro Ala Glu Ala Trp Asn
145 150 155 160

30 Phe Trp Thr Arg Lys Gly Leu Val Ser Gly Gly Leu Tyr Glu Ser His
165 170 175

35 Val Gly Cys Arg Pro Tyr Ser Ile Pro Pro Cys Glu His His Val Asn
180 185 190

40 Gly Ser Arg Pro Pro Cys Thr Gly Glu Gly Asp Thr Pro Lys Cys Ser
195 200 205

45 Lys Ile Cys Glu Pro Gly Tyr Ser Pro Thr Tyr Lys Gln Asp Lys His
210 215 220

50 Tyr Gly Tyr Asn Ser Tyr Ser Val Ser Asn Ser Glu Lys Asp Ile Met
225 230 235 240

55 Ala Glu Ile Tyr Lys Asn Gly Pro Val Glu Gly Ala Phe Ser Val Tyr
245 250 255

60 Ser Asp Phe Leu Leu Tyr Lys Ser Gly Val Tyr Gln His Val Thr Gly
260 265 270

65 Glu Met Met Gly Gly His Ala Ile Arg Ile Leu Gly Trp Gly Val Glu
275 280 285

70 Asn Gly Thr Pro Tyr Trp Leu Val Ala Asn Ser Trp Asn Thr Asp Trp
290 295 300

75 Gly Asp Asn Gly Phe Phe Lys Ile Leu Arg Gly Gln Asp His Cys Gly
305 310 315 320

80 Ile Glu Ser Glu Val Val Ala Gly Ile Pro Arg Thr Asp Gln Tyr Trp
325 330 335

Glu Lys Ile

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 10 <300>
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 <313> (1)..(1249)
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 15 Met Ala Thr Ala Ala Thr Glu Glu Pro Phe Pro Phe His Gly Leu Leu
 1 5 10 15
 20 Pro Lys Lys Glu Thr Gly Ala Ala Ser Phe Leu Cys Arg Tyr Pro Glu
 20 25 30
 25 Tyr Asp Gly Arg Gly Val Leu Ile Ala Val Leu Asp Thr Gly Val Asp
 35 40 45
 30 Pro Gly Ala Pro Gly Met Gln Val Thr Thr Asp Gly Lys Pro Lys Ile
 50 55 60
 35 Val Asp Ile Ile Asp Thr Thr Gly Ser Gly Asp Val Asn Thr Ala Thr
 65 70 75 80
 40 Glu Val Glu Pro Lys Asp Gly Glu Ile Val Gly Leu Ser Gly Arg Val
 85 90 95
 45 Leu Lys Ile Pro Ala Ser Trp Thr Asn Pro Ser Gly Lys Tyr His Ile
 100 105 110
 50 Gly Ile Lys Asn Gly Tyr Asp Phe Tyr Pro Lys Ala Leu Lys Glu Arg
 115 120 125
 55 Ile Gln Lys Glu Arg Lys Glu Lys Ile Trp Asp Pro Val His Arg Val
 130 135 140
 60 Ala Leu Ala Glu Ala Cys Arg Lys Gln Glu Glu Phe Asp Val Ala Asn
 145 150 155 160
 65 Asn Gly Ser Ser Gln Ala Asn Lys Leu Ile Lys Glu Glu Leu Gln Ser
 165 170 175
 70 Gln Val Glu Leu Leu Asn Ser Phe Glu Lys Lys Tyr Ser Asp Pro Gly
 180 185 190
 75 Pro Val Tyr Asp Cys Leu Val Trp His Asp Gly Glu Val Trp Arg Ala
 195 200 205
 80 Cys Ile Asp Ser Asn Glu Asp Gly Asp Leu Ser Lys Ser Thr Val Leu
 210 215 220
 85 Arg Asn Tyr Lys Glu Ala Gln Glu Tyr Gly Ser Phe Gly Thr Ala Glu

95

225 230 235 240

5 Met Leu Asn Tyr Ser Val Asn Ile Tyr Asp Asp Gly Asn Leu Leu Ser
245 250 25510 Ile Val Thr Ser Gly Gly Ala His Gly Thr His Val Ala Ser Ile Ala
260 265 270Ala Gly His Phe Pro Glu Glu Pro Glu Arg Asn Gly Val Ala Pro Gly
275 280 28515 Ala Gln Ile Leu Ser Ile Lys Ile Gly Asp Thr Arg Leu Ser Thr Met
290 295 30020 Glu Thr Gly Thr Gly Leu Ile Arg Ala Met Ile Glu Val Ile Asn His
305 310 315 32025 Lys Cys Asp Leu Val Asn Tyr Ser Tyr Gly Glu Ala Thr His Trp Pro
325 330 33530 Asn Ser Gly Arg Ile Cys Glu Val Ile Asn Glu Ala Val Trp Lys His
340 345 35035 Asn Ile Ile Tyr Val Ser Ser Ala Gly Asn Asn Gly Pro Cys Leu Ser
355 360 365

370 Gly Cys Pro Gly Gly Thr Thr Ser Ser Val Ile Gly Val Gly

40 Ala Tyr Val Ser Pro Asp Met Met Val Ala Glu Tyr Ser Leu Arg Glu
385 390 395 40045 Lys Leu Pro Ala Asn Gln Tyr Thr Trp Ser Ser Arg Gly Pro Ser Ala
405 410 41550 Asp Gly Ala Leu Gly Val Ser Ile Ser Ala Pro Gly Gly Ala Ile Ala
420 425 43055 Ser Val Pro Asn Trp Thr Leu Arg Gly Thr Gln Leu Met Asn Gly Thr
435 440 44560 Ser Met Ser Ser Pro Asn Ala Cys Gly Gly Ile Ala Leu Ile Leu Ser
450 455 46065 Gly Leu Lys Ala Asn Asn Ile Asp Tyr Thr Val His Ser Val Arg Arg
465 470 475 48070 Ala Leu Glu Asn Thr Ala Val Lys Ala Asp Asn Ile Glu Val Phe Ala
485 490 495Gln Gly His Gly Ile Ile Gln Val Asp Lys Ala Tyr Asp Tyr Leu Val
500 505 510

Gln Asn Thr Ser Phe Ala Asn Lys Leu Gly Phe Thr Val Thr Val Gly
515 520 525

5 Asn Asn Arg Gly Ile Tyr Leu Arg Asp Pro Val Gln Val Ala Ala Pro
530 535 540

10 Ser Asp His Gly Val Gly Ile Glu Pro Val Phe Pro Glu Asn Thr Glu
545 550 555 560

15 Asn Ser Glu Lys Ile Ser Leu Gln Leu His Leu Ala Leu Thr Ser Asn
565 570 575

20 Ser Ser Trp Val Gln Cys Pro Ser His Leu Glu Leu Met Asn Gln Cys
580 585 590

Arg His Ile Asn Ile Arg Val Asp Pro Arg Gly Leu Arg Glu Gly Leu
595 600 605

25 His Tyr Thr Glu Val Cys Gly Tyr Asp Ile Ala Ser Pro Asn Ala Gly
610 615 620

30 Pro Leu Phe Arg Val Pro Ile Thr Ala Val Ile Ala Ala Lys Val Asn
625 630 635 640

35 Glu Ser Ser His Tyr Asp Leu Ala Phe Thr Asp Val His Phe Lys Pro
645 650 655

40 Gly Gln Ile Arg Arg His Phe Ile Glu Val Pro Glu Gly Ala Thr Trp
660 665 670

Ala Glu Val Thr Val Cys Ser Cys Ser Ser Glu Val Ser Ala Lys Phe
675 680 685

45 Val Leu His Ala Val Gln Leu Val Lys Gln Arg Ala Tyr Arg Ser His
690 695 700

50 Glu Phe Tyr Lys Phe Cys Ser Leu Pro Glu Lys Gly Thr Leu Thr Glu
705 710 715 720

55 Ala Phe Pro Val Leu Gly Gly Lys Ala Ile Glu Phe Cys Ile Ala Arg
725 730 735

Trp Trp Ala Ser Leu Ser Asp Val Asn Ile Asp Tyr Thr Ile Ser Phe
740 745 750

60 His Gly Ile Val Cys Thr Ala Pro Gln Leu Asn Ile His Ala Ser Glu
755 760 765

65 Gly Ile Asn Arg Phe Asp Val Gln Ser Ser Leu Lys Tyr Glu Asp Leu
770 775 780

70 Ala Pro Cys Ile Thr Leu Lys Asn Trp Val Gln Thr Leu Arg Pro Val

	785	790	795	800
	...			
5	Ser Ala Lys Thr Lys Pro Leu Gly Ser Arg Asp Val Leu Pro Asn Asn 805 810 815			
10	Arg Gln Leu Tyr Glu Met Val Leu Thr Tyr Asn Phe His Gln Pro Lys 820 825 830			
15	Ser Gly Glu Val Thr Pro Ser Cys Pro Leu Leu Cys Glu Leu Leu Tyr 835 840 845			
20	Glu Ser Glu Phe Asp Ser Gln Leu Trp Ile Ile Phe Asp Gln Asn Lys 850 855 860			
25	Arg Gln Met Gly Ser Gly Asp Ala Tyr Pro His Gln Tyr Ser Leu Lys 865 870 875 880			
30	Leu Glu Lys Gly Asp Tyr Thr Ile Arg Leu Gln Ile Arg His Glu Gln 885 890 895			
35	Ile Ser Asp Leu Glu Arg Leu Lys Asp Leu Pro Phe Ile Val Ser His 900 905 910			
40	Arg Leu Ser Asn Thr Leu Ser Leu Asp Ile His Glu Asn His Ser Phe 915 920 925			
45	Ala Leu Leu Gly Lys Lys Ser Ser Asn Leu Thr Leu Pro Pro Lys 930 935 940			
50	Tyr Asn Gln Pro Phe Phe Val Thr Ser Leu Pro Asp Asp Lys Ile Pro 945 950 955 960			
55	Lys Gly Ala Gly Pro Gly Cys Tyr Leu Ala Gly Ser Leu Thr Leu Ser 965 970 975			
60	Lys Thr Glu Leu Gly Lys Lys Ala Asp Val Ile Pro Val His Tyr Tyr 980 985 990			
65	Leu Ile Pro Pro Pro Thr Lys Thr Lys Asn Gly Ser Lys Asp Lys Glu 995 1000 1005			
70	Lys Asp Ser Glu Lys Glu Lys Asp Leu Lys Glu Glu Phe Thr Glu 1010 1015 1020			
75	Ala Leu Arg Asp Leu Lys Ile Gln Trp Met Thr Lys Leu Asp Ser 1025 1030 1035			
80	Ser Asp Ile Tyr Asn Glu Leu Lys Glu Thr Tyr Pro Asn Tyr Leu 1040 1045 1050			
85	Pro Leu Tyr Val Ala Arg Leu His Gln Leu Asp Ala Glu Lys Glu 1055 1060 1065			

Arg Met Lys Arg Leu Asn Glu Ile Val Asp Ala Ala Asn Ala Val
 1070 1075 1080
 5 Ile Ser His Ile Asp Gln Thr Ala Leu Ala Val Tyr Ile Ala Met
 1085 1090 1095
 10 Lys Thr Asp Pro Arg Pro Asp Ala Ala Thr Ile Lys Asn Asp Met
 1100 1105 1110
 15 Asp Lys Gln Lys Ser Thr Leu Val Asp Ala Leu Cys Arg Lys Gly
 1115 1120 1125
 20 Cys Ala Leu Ala Asp His Leu Leu His Thr Gln Ala Gln Asp Gly
 1130 1135 1140
 25 Ala Ile Ser Thr Asp Ala Glu Gly Lys Glu Glu Glu Gly Glu Ser
 1145 1150 1155
 30 Pro Leu Asp Ser Leu Ala Glu Thr Phe Trp Glu Thr Thr Lys Trp
 1160 1165 1170
 35 Thr Asp Leu Phe Asp Asn Lys Val Leu Thr Phe Ala Tyr Lys His
 1175 1180 1185
 40 Ala Leu Val Asn Lys Met Tyr Gly Arg Gly Leu Lys Phe Ala Thr
 1190 1195 1200
 Lys Leu Val Glu Glu Lys Pro Thr Lys Glu Asn Trp Lys Asn Cys
 1205 1210 1215
 45 Ile Gln Leu Met Lys Leu Leu Gly Trp Thr His Cys Ala Ser Phe
 1220 1225 1230
 Thr Glu Asn Trp Leu Pro Ile Met Tyr Pro Pro Asp Tyr Cys Val
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 50 Phe
 55 <210> 138
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 <309> 1997-11-01
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 65 <400> 138
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 70 Ala Val Pro Ile Asp Asp Pro Glu Asp Gly Gly Lys His Trp Val Val

99

20

25

30

5 Ile Val Ala Gly Ser Asn Gly Trp Tyr Asn Tyr Arg His Gln Ala Asp
 35 40 45
 10 Ala Cys His Ala Tyr Gln Ile Ile His Arg Asn Gly Ile Pro Asp Glu
 50 55 60
 15 Gln Ile Val Val Met Met Tyr Asp Asp Ile Ala Tyr Ser Glu Asp Asn
 65 70 75 80
 20 Pro Thr Pro Gly Ile Val Ile Asn Arg Pro Asn Gly Thr Asp Val Tyr
 85 90 95
 25 Gln Gly Val Pro Lys Asp Tyr Thr Gly Glu Asp Val Thr Pro Gln Asn
 100 105 110
 30 Phe Leu Ala Val Leu Arg Gly Asp Ala Glu Ala Val Lys Gly Ile Gly
 115 120 125
 35 Ser Gly Lys Val Leu Lys Ser Gly Pro Gln Asp His Val Phe Ile Tyr
 130 135 140
 40 Phe Thr Asp His Gly Ser Thr Gly Ile Leu Val Phe Pro Asn Glu Asp
 145 150 155 160
 45 Leu His Val Lys Asp Leu Asn Glu Thr Ile His Tyr Met Tyr Lys His
 165 170 175
 50 Lys Met Tyr Arg Lys Met Val Phe Tyr Ile Glu Ala Cys Glu Ser Gly
 180 185 190
 55 Ser Met Met Asn His Leu Pro Asp Asn Ile Asn Val Tyr Ala Thr Thr
 195 200 205
 60 Ala Ala Asn Pro Arg Glu Ser Ser Tyr Ala Cys Tyr Tyr Asp Glu Lys
 210 215 220
 65 Arg Ser Thr Tyr Leu Gly Asp Trp Tyr Ser Val Asn Trp Met Glu Asp
 225 230 235 240
 70 Ser Asp Val Glu Asp Leu Thr Lys Glu Thr Leu His Lys Gln Tyr His
 245 250 255
 75 Leu Val Lys Ser His Thr Asn Thr Ser His Val Met Gln Tyr Gly Asn
 260 265 270
 80 Lys Thr Ile Ser Thr Met Lys Val Met Gln Phe Gln Gly Met Lys Arg
 275 280 285
 85 Lys Ala Ser Ser Pro Val Pro Leu Pro Pro Val Thr His Leu Asp Leu
 290 295 300

100

Thr Pro Ser Pro Asp Val Pro Leu Thr Ile Met Lys Arg Lys Leu Met
305 310 315 320

5 Asn Thr Asn Asp Leu Glu Glu Ser Arg Gln Leu Thr Glu Glu Ile Gln
325 330 335

10 Arg His Leu Asp Ala Arg His Leu Ile Glu Lys Ser Val Arg Lys Ile
340 345 350

15 Val Ser Leu Leu Ala Ala Ser Glu Ala Glu Val Glu Gln Leu Leu Ser
355 360 365

20 Glu Arg Ala Pro Leu Thr Gly His Ser Cys Tyr Pro Glu Ala Leu Leu
370 375 380

25 His Phe Arg Thr His Cys Phe Asn Trp His Ser Pro Thr Tyr Glu Tyr
385 390 395 400

30 Ala Leu Arg His Leu Tyr Val Leu Val Asn Leu Cys Glu Lys Pro Tyr
405 410 415

35 Tyr

35 <210> 139

40 <211> 342

<212> PRT

<213> Homo sapiens

45 <300>

<308> Swissprot/P25105

<309> 1992-05-01

<313> (1)..(342)

50 <400> 139

Met Glu Pro His Asp Ser Ser His Met Asp Ser Glu Phe Arg Tyr Thr
1 5 10 15

55 Leu Phe Pro Ile Val Tyr Ser Ile Ile Phe Val Leu Gly Val Ile Ala
20 25 30

60 Asn Gly Tyr Val Leu Trp Val Phe Ala Arg Leu Tyr Pro Cys Lys Lys
35 40 45

65 Phe Asn Glu Ile Lys Ile Phe Met Val Asn Leu Thr Met Ala Asp Met
50 55 60

Leu Phe Leu Ile Thr Leu Pro Leu Trp Ile Val Tyr Tyr Gln Asn Gln
65 70 75 80

70 Gly Asn Trp Ile Leu Pro Lys Phe Leu Cys Asn Val Ala Gly Cys Leu

101

85

90

95

5 Phe Phe Ile Asn Thr Tyr Cys Ser Val Ala Phe Leu Gly Val Ile Thr
 100 105 110
 10 Tyr Asn Arg Phe Gln Ala Val Thr Arg Pro Ile Lys Thr Ala Gln Ala
 115 120 125
 15 Asn Thr Arg Lys Arg Gly Ile Ser Leu Ser Leu Val Ile Trp Val Ala
 130 135 140
 20 Ile Val Gly Ala Ala Ser Tyr Phe Leu Ile Leu Asp Ser Thr Asn Thr
 145 150 155 160
 25 Val Pro Asp Ser Ala Gly Ser Gly Asn Val Thr Arg Cys Phe Glu His
 165 170 175
 30 Tyr Glu Lys Gly Ser Val Pro Val Leu Ile Ile His Ile Phe Ile Val
 180 185 190
 35 Phe Ser Phe Phe Leu Val Phe Leu Ile Ile Leu Phe Cys Asn Leu Val
 195 200 205
 40 Ile Ile Arg Thr Leu Leu Met Gln Pro Val Gln Gln Gln Arg Asn Ala
 210 215 220
 45 Glu Val Lys Arg Arg Ala Leu Trp Met Val Cys Thr Val Leu Ala Val
 225 230 235 240
 50 Phe Ile Ile Cys Phe Val Pro His His Val Val Gln Leu Pro Trp Thr
 245 250 255
 55 Leu Ala Glu Leu Gly Phe Gln Asp Ser Lys Phe His Gln Ala Ile Asn
 260 265 270
 60 Asp Ala His Gln Val Thr Leu Cys Leu Leu Ser Thr Asn Cys Val Leu
 275 280 285
 65 Asp Pro Val Ile Tyr Cys Phe Leu Thr Lys Lys Phe Arg Lys His Leu
 290 295 300
 70 Thr Glu Lys Phe Tyr Ser Met Arg Ser Ser Arg Lys Cys Ser Arg Ala
 305 310 315 320
 75 Thr Thr Asp Thr Val Thr Glu Val Val Val Pro Phe Asn Gln Ile Pro
 325 330 335
 80 Gly Asn Ser Leu Lys Asn
 340
 <210> 140
 <211> 359
 <212> PRT

<213> Homo sapiens

<300>

<308> Swissprot/Q92187

5 <309> 1997-11-01

<313> (1)..(359)

<400> 140

10 Met Arg Ser Ile Arg Lys Arg Trp Thr Ile Cys Thr Ile Ser Leu Leu
1 5 10 15

15 Leu Ile Phe Tyr Lys Thr Lys Glu Ile Ala Arg Thr Glu Glu His Gln
20 25 30

20 Glu Thr Gln Leu Ile Gly Asp Gly Glu Leu Ser Leu Ser Arg Ser Leu
35 40 45

25 Val Asn Ser Ser Asp Lys Ile Ile Arg Lys Ala Gly Ser Ser Ile Phe
50 55 60

30 Gln His Asn Val Glu Gly Trp Lys Ile Asn Ser Ser Leu Val Leu Glu
65 70 75 80

35 Ile Arg Lys Asn Ile Leu Arg Phe Leu Asp Ala Glu Arg Asp Val Ser
85 90 95

40 Val Val Lys Ser Ser Phe Lys Pro Gly Asp Val Ile His Tyr Val Leu
100 105 110

45 Asp Arg Arg Arg Thr Leu Asn Ile Ser His Asp Leu His Ser Leu Leu
115 120 125

50 Pro Glu Val Ser Pro Met Lys Asn Arg Arg Phe Lys Thr Cys Ala Val
130 135 140

55 Val Gly Asn Ser Gly Ile Leu Leu Asp Ser Glu Cys Gly Lys Glu Ile
145 150 155 160

60 Asp Ser His Asn Phe Val Ile Arg Cys Asn Leu Ala Pro Val Val Glu
165 170 175

65 Phe Ala Ala Asp Val Gly Thr Lys Ser Asp Phe Ile Thr Met Asn Pro
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70 Ser Val Val Gln Arg Ala Phe Gly Gly Phe Arg Asn Glu Ser Asp Arg
195 200 205

75 Glu Lys Phe Val His Arg Leu Ser Met Leu Asn Asp Ser Val Leu Trp
210 215 220

80 Ile Pro Ala Phe Met Val Lys Gly Gly Glu Lys His Val Glu Trp Val
225 230 235 240

85 Asn Ala Leu Ile Leu Lys Asn Lys Leu Lys Val Arg Thr Ala Tyr Pro

103

245

250

255

5 Ser Leu Arg Leu Ile His Ala Val Arg Gly Tyr Trp Leu Thr Asn Lys
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 10 Val Pro Ile Lys Arg Pro Ser Thr Gly Leu Leu Met Tyr Thr Leu Ala
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 15 Thr Arg Phe Cys Asp Glu Ile His Leu Tyr Gly Phe Trp Pro Phe Pro
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 20 Lys Asp Leu Asn Gly Lys Ala Val Lys Tyr His Tyr Tyr Asp Asp Leu
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 25 Lys Tyr Arg Tyr Phe Ser Asn Ala Ser Pro His Arg Met Pro Leu Glu
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 35 40 45
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 65 Trp Asp Thr Ala Gly Gln Glu Arg Tyr Arg Ala Ile Thr Ser Ala Tyr
 70 75 80
 75 Tyr Arg Gly Ala Val Gly Ala Leu Leu Val Tyr Asp Ile Ala Lys His
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 80 Leu Thr Tyr Glu Asn Val Glu Arg Trp Leu Lys Glu Leu Arg Asp His
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Ala Asp Ser Asn Ile Val Ile Met Leu Val Gly Asn Lys Ser Asp Leu
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5 Arg His Leu Arg Ala Val Pro Thr Asp Glu Ala Arg Ala Phe Ala Glu
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10 Lys Asn Asn Leu Ser Phe Ile Glu Thr Ser Ala Leu Asp Ser Thr Asn
145 150 155 160

15 Val Glu Glu Ala Phe Lys Asn Ile Leu Thr Glu Ile Tyr Arg Ile Val
165 170 175

20 Ser Gln Lys Gln Ile Ala Asp Arg Ala Ala His Asp Glu Ser Pro Gly
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Asn Asn Val Val Asp Ile Ser Val Pro Pro Thr Thr Asp Gly Gln Lys
195 200 205

25 Pro Asn Lys Leu Gln Cys Cys Gln Asn Leu
210 215

30